

# FIGURE 1

| 10           | 20          | 30          | 40          | 50         |     |
|--------------|-------------|-------------|-------------|------------|-----|
| 1234567890   | 1234567890  | 1234567890  | 1234567890  | 1234567890 |     |
| TGGCCACTOC   | CTCTCTGCGC  | GCTCGCTCGC  | TCACTGAGGC  | GGGGCGACCA | 50  |
| AAGGTGCGCC   | GACGCCCGGG  | CITTCGCCCGG | GAGGCCTCAG  | TGAGCGAGCG | 100 |
| AGCGCGCAGA   | GAGGGAGTGG  | CCAACCTCCAT | CACTAGGGT   | TCCTCAGATC | 150 |
| TCTTCTAAG    | TAAACAGTAC  | ATGAACCTTT  | ACCCCGTTGC  | TGGCAACCGG | 200 |
| OCTGGTCTGT   | GCCAAGTGT   | TGCTGACGCA  | ACCCCCACTG  | GCTGGGGCTT | 250 |
| GGCCATAGGC   | CATCAGCGCA  | TGCGGATCTC  | AGTGTGGTTT  | TGCAAGAGGA | 300 |
| AGCAAAAAGC   | CCTCTCCACCC | AGGCCTGGAA  | TGTTCCACC   | CAATGTCGAG | 350 |
| CAGTGTGGTT   | TIGCAAGAGG  | AAGCAAAAAG  | CCTCTCCACC  | CAGGCCTGGA | 400 |
| CTCGAGAGCT   | TGACCCACCA  | TGCAAATAGA  | GCTCTCCACC  | TGCTTCTTTC | 450 |
| M etGlnIleG1 | uLeuSerThr  | CysPhePheL  |             |            |     |
| TGTGCCTTT    | GCGATTCTGC  | TTTAGTGCCA  | CCAGAAGATA  | CTACCTGGGT | 500 |
| euCysLeuLe   | uArgPheCys  | PheSerAlaT  | hrArgArgTy  | rTyrLeuGly |     |
| GCAGTGGAAC   | TGTATGGGA   | CTATATGCAA  | AGTGATCTCG  | GTGAGCTGCC | 550 |
| AlaValGluL   | euSerTrpAs  | pTyrMetGln  | SerAspLeuG  | lyGluLeuPr |     |
| TGTGGACGCA   | AGATTCTCTC  | CTAGAGTGCC  | AAAATCTTT   | CCATTCAACA | 600 |
| oValAspAla   | ArgPheProP  | roArgValPr  | oLysSerPhe  | ProPheAsnT |     |
| CCTCAGTCGT   | GTACAAAAAG  | ACTCTGTTTG  | TAGAATTICAC | GGITCACCTT | 650 |
| hrSerValVa   | 1TyrLysLys  | ThrLeuPheV  | alGluPheTh  | rValHisLeu |     |
| TTCAACATCG   | CTAACCCAAG  | GOCAACCTGG  | ATGGGTCTGC  | TAGGTCTTAC | 700 |
| PheAsnIleA   | 1aLysProAr  | gProProTrp  | MetGlyLeuL  | euGlyProTh |     |
| CATCCAGGCT   | GAGGTTATG   | ATACAGTGGT  | CATTACACIT  | AAGAACATGG | 750 |
| rIleGlnAla   | GluValTyrA  | spThrValVa  | 1IleThrLeu  | LysAsnMetA |     |
| CTTOCCATOC   | TGTCAGCTT   | CATGCTGTTG  | GTGTATCTA   | CTGGAAAGCT | 800 |
| 1aSerHisPr   | oValSerLeu  | HisAlaValG  | 1yValSerTy  | rTrpLysAla |     |
| TCTGAGGGAG   | CTGAATATGA  | TGATCAGACCC | AGTCAGGAGG  | AGAAAGAAGA | 850 |
| SerGluGlyA   | 1aGluTyrAs  | pAspGlnThr  | SerGlnArgG  | 1uLysGluAs |     |
| TGATAAAAGTC  | TTCCCTGGTG  | GAAGCCATAC  | ATATGCTGG   | CAGGTCTTGA | 900 |
| pAspLysVal   | PheProGlyG  | lySerHisTh  | rTyrValTrp  | GlnValLeuL |     |
| AAGAGAATGG   | TCCAATGGCC  | TCTGACCCAC  | TGTGCCTTAC  | CTACTCATAT | 950 |
| ysGluAsnG1   | yProMetAla  | SerAspProL  | euCysLeuTh  | rTyrSerTyr |     |

FIGURE 1 (cont.)

| 10         | 20          | 30         | 40          | 50          |      |
|------------|-------------|------------|-------------|-------------|------|
| 1234567890 | 1234567890  | 1234567890 | 1234567890  | 1234567890  |      |
| CTTCTCATG  | TGGACCTGGT  | AAAAGACTTG | AATTCAAGGCC | TCATTGGAGC  | 1000 |
| LeuSerHisV | aLAspLeuVa  | lLysAspLeu | AsnSerGlyL  | euIleGlyAl  |      |
| CCTACTAGTA | TGTAGAGAAG  | GGAGTCTGGC | CAAGGAAAAG  | ACACAGACCT  | 1050 |
| aLeuLeuVal | CysArgGluG  | lySerLeuAl | aLysGluLys  | ThrGlnThrL  |      |
| TGCACAAATT | TATACTACTT  | TTTGCIGTAT | TTGATGAAGG  | GAAAAGTGG   | 1100 |
| euHisLysPh | eIleLeuLeu  | PheAlaValP | heAspGluG1  | yLysSerTrp  |      |
| CACICAGAAA | CAAAGAACTC  | CTTGATGCAG | GATAGGGATG  | CTGCATCTGC  | 1150 |
| HisSerGluT | hrLysAsnSe  | rLeuMetGln | AspArgAspA  | laAlaSerAl  |      |
| TOGGGCTGG  | CCTAAAATGC  | ACACAGTCAA | TGGTTATGTA  | AACAGGTCTC  | 1200 |
| aArgAlaTrp | ProLysMetH  | isThrValAs | nGlyTyrVal  | AsnArgSerL  |      |
| TGCCAGGTCT | GATTGGATGC  | CACAGGAAAT | CAGTCTATTG  | GCATGTGATT  | 1250 |
| euProGlyLe | uIleGlyCys  | HisArgLysS | erValTyrTr  | pHisValIle  |      |
| GGAATGGGCA | CCACTCCTGA  | AGTGCACICA | ATATTCCCTCG | AAGGTACACAC | 1300 |
| GlyMetGlyT | hrThrProG1  | uValHisSer | IlePheLeuG  | luGlyHisTh  |      |
| ATTCTTGIG  | AGGAACCATC  | GCCAGGCGTC | CTTGGAAATC  | TOGCCAATAA  | 1350 |
| rPheLeuVal | ArgAsnHisA  | rgGlnAlaSe | rLeuGluIle  | SerProIleT  |      |
| CTTCCCTTAC | TGCTCAAACA  | CTCTTGATGG | ACCTTGGACA  | TTTTCTACTG  | 1400 |
| hrPheLeuTh | rAlaGlnThr  | LeuLeuMetA | spLeuGlyG1  | nPheLeuLeu  |      |
| TTTGTICATA | TCTCTTCCCA  | CCAACATGAT | GGCATGGAAG  | CTTATGICAA  | 1450 |
| PheCysHisI | leSerSerHi  | sGlnHisAsp | GlyMetGluA  | laTyrValLy  |      |
| AGTAGACAGC | TGTCCAGAGG  | AACCCCAACT | ACGAATGAAA  | AATAATGAAG  | 1500 |
| sValAspSer | CysProGluG  | luProGlnLe | uArgMetLys  | AsnAsnGluG  |      |
| AAGCGGAAGA | CTATGATGAT  | GATCTTACTG | ATTCTGAAAT  | GGATGTGGTC  | 1550 |
| luAlaGluAs | pTyrAspAsp  | AspLeuThrA | spSerGluMe  | tAspValVal  |      |
| AGGTTTGATG | ATGACAACTC  | TCCTCCCTT  | ATCCAAATTG  | GCTCAGTTC   | 1600 |
| ArgPheAspA | spAspAsnSe  | rProSerPhe | IleGlnIleA  | rgSerValAl  |      |
| CAAGAAGCAT | CCTAAAACCTT | GGGTACATTA | CATTGCTGCT  | GAAGAGGAGG  | 1650 |
| aLysLysHis | ProLysThrT  | rpValHistY | rIleAlaAla  | GluGluGluA  |      |
| ACTGGGACTA | TGCTCCCTTA  | GTCCTCGGCC | CCGATGACAG  | AAGTATAAAA  | 1700 |
| spTrpAspTy | rAlaProLeu  | ValLeuAlaP | roAspAspAr  | gSerTyrLys  |      |
| AGTCAATATT | TGAACAATGG  | CCCTCAGCGG | ATTGGTAGGA  | AGTACAAAAAA | 1750 |
| SerGlnTyrL | euAsnAsnG1  | yProGlnArg | IleGlyArgL  | ysTyrLysLy  |      |
| AGTCCGATT  | ATGGCATAACA | CAGATGAAAC | CTTTAAGACT  | CGTGAAGCTA  | 1800 |
| sValArgPhe | MetAlaTyrT  | hrAspGluTh | rPheLysThr  | ArgGluAlaI  |      |
| TTCAGCATGA | ATCAGGAATC  | TTGGGACCTT | TACTTTATGG  | GGAAGTGG    | 1850 |
| leGlnHisG1 | uSerGlyIle  | LeuGlyProL | euLeuTyrG1  | yGluValGly  |      |
| GACACACTGT | TGATTATATT  | TAAGAATCAA | GCAAGCAGAC  | CATATAACAT  | 1900 |
| AspThrLeuL | euIleIlePh  | eLysAsnGln | AlaSerArgP  | roTyrAsnII  |      |

FIGURE 1 (cont.)

| 10           | 20         | 30          | 40         | 50          |      |
|--------------|------------|-------------|------------|-------------|------|
| 1234567890   | 1234567890 | 1234567890  | 1234567890 | 1234567890  |      |
| CTACCCCTCAC  | GGAATCACTG | ATGTOCGTCC  | TTTGTATTCA | AGGAGATTAC  | 1950 |
| eTyrProHis   | GlyIleThrA | spValArgPr  | oLeuTyrSer | ArgArgLeuP  |      |
| CAAAAGGTGT   | AAAACATTIG | AAGGATTTTC  | CAATTCTGCC | AGGAGAAATA  | 2000 |
| roLysGlyVa   | 1LysHisLeu | LysAspPheP  | roIleLeuPr | oGlyGluIle  |      |
| TTCAAATATA   | AATGGACAGT | GACTGTAGAA  | GATGGGCCAA | CTAAATCAGA  | 2050 |
| PheLysTyrL   | ysTrpThrVa | 1ThrValGlu  | AspGlyProT | hrLysSerAs  |      |
| TCCTCGGTGC   | CTGACCCGCT | ATTACTCTAG  | TTTCGTTAAT | ATGGAGAGAG  | 2100 |
| pProArgCys   | LeuThrArgT | yrTyrSerSe  | rPheValAsn | MetGluArgA  |      |
| ATCTAGCTTC   | AGGACTCATT | GGCCCTCTCC  | TCATCTGCTA | CAAAGAACCT  | 2150 |
| spLeuAlaSe   | rGlyLeuIle | GlyProLeuL  | euIleCysTy | rLysGluSer  |      |
| GTAGATCAAA   | GAGGAAACCA | GATAATGTCA  | GACAAGAGGA | ATGTCATCCT  | 2200 |
| ValAspGlnA   | rgGlyAsnG1 | nIleMetSer  | AspLysArgA | snValIleLe  |      |
| GTTTCTGTCA   | TTTGATGAGA | ACCGAAGCTG  | GTACCTCACA | GAGAATATAC  | 2250 |
| uPheSerVal   | PheAspGluA | snArgSerTr  | pTyrLeuThr | GluAsnIleG  |      |
| AACGCTTTCT   | CCCCAATCCA | GCTGGAGTGC  | AGCTTGAGGA | TOCAGAGTTC  | 2300 |
| 1nArgPheLe   | uProAsnPro | AlaGlyValG  | 1nLeuGluAs | pProGluPhe  |      |
| CAAGOCTCCA   | ACATCATGCA | CAGCATCAAT  | GGCTATGTTT | TTGATAGTTT  | 2350 |
| GlnAlaSerA   | snIleMetHi | sSerIleAsn  | GlyTyrValP | heAspSerLe  |      |
| GCAGTTGTCA   | GTTGTTTGC  | ATGAGGTGGC  | ATACTGGTAC | ATTCTAACGA  | 2400 |
| uGlnLeuSer   | ValCysLeuH | isGluValAl  | aTyrTrpTyr | IleLeuSerI  |      |
| TTGGAGCACCA  | GACTGACTTC | CTTCTGTCT   | TCTTCTCTGG | ATATAACCTTC | 2450 |
| 1eGlyAlaG1   | nThrAspPhe | LeuSerValP  | hePheSerG1 | yTyrThrPhe  |      |
| AAACACAAAA   | TGGTCTATGA | AGACACACTC  | ACCCATTCTC | CATTCTCAGG  | 2500 |
| LysHisLysM   | etValTyrG1 | uAspThrLeu  | ThrLeuPheP | roPheSerG1  |      |
| AGAAAACGTGTC | TTCATGTGCA | TGGAAAACCC  | AGGICTATGG | ATTCTGGGGT  | 2550 |
| yGluThrVal   | PheMetSerM | etGluAsnPr  | oGlyLeuTrp | IleLeuGlyC  |      |
| GCCACAACTC   | AGACTTTGG  | AACAGAGGCCA | TGACCGCCTT | ACTGAAGGTT  | 2600 |
| ysHisAsnSe   | rAspPheArg | AsnArgGlyM  | etThrAlaLe | uLeuLysVal  |      |
| TCTAGTTGTG   | ACAAGAACAC | TGGTGATTAT  | TACGAGGACA | GTTATGAAGA  | 2650 |
| SerSerCysA   | spLysAsnTh | rGlyAspTyr  | TyrGluAspS | erTyrGluAs  |      |
| TATTTCAGCA   | TACTTGCTGA | GTAAAAACAA  | TGCCATTGAA | CCAAGAACCT  | 2700 |
| pIleSerAla   | TyrLeuLeuS | erLysAsnAs  | nAlaIleGlu | ProArgSerP  |      |
| TCTCCAGAA    | TICAAGACAC | CCTAGCACTA  | GGCAAAAGCA | ATTTAATGCC  | 2750 |
| heSerGlnAs   | nSerArgHis | ProSerThrA  | rgGlnLysG1 | nPheAsnAla  |      |
| ACCCCACCAAG  | TCTTGAAACG | CCATCAACGG  | GAAATAACTC | GTACTACTCT  | 2800 |
| ThrProProV   | alLeuLysAr | gHisGlnArg  | GluIleThrA | rgThrThrLe  |      |
| TCAGTCAGAT   | CAAGAGGAAA | TTGACTATGA  | TGATACCATA | TCAGITGAAA  | 2850 |
| uGlnSerAsp   | GlnGluGluI | leAspTyrAs  | pAspThrIle | SerValGluM  |      |

FIGURE 1 (cont.)

|            | 10          | 20         | 30         | 40          | 50         |      |
|------------|-------------|------------|------------|-------------|------------|------|
| 1234567890 | 1234567890  | 1234567890 | 1234567890 | 1234567890  | 1234567890 | 2900 |
| TGAAGAAGGA | AGATTTGAC   | ATTATGATG  | AGGATGAAAA | TCAGAGCCCC  |            |      |
| etLysLysG1 | uAspPheAsp  | IleTyrAspG | luAspGluAs | nGlnSerPro  |            |      |
| CGCAGCTTTC | AAAAGAAAAC  | ACGACACTAT | TTTATTGCTG | CAGTGGAGAG  |            |      |
| ArgSerPheG | InLysLysTh  | rArgHisTyr | PheIleAlaA | laValGluAr  |            |      |
| gLeuTrpAsp | TATGGGATGA  | GTAGCTCCCC | ACATGTTCTA | AGAAACAGGG  |            |      |
| CTCAGAGTGG | CAGTGTCCCT  | CAGTCAAGA  | AAGTGTGTTT | CCAGGAATT   |            |      |
| laGlnSerG1 | ySerValPro  | GlnPheLysL | ysValValPh | eGlnGluPhe  |            |      |
| ACTGATGGCT | CCTTTACTCA  | GCCCTTATAC | CGTGGAGAAC | TAAATGAACA  |            |      |
| ThrAspGlyS | erPheThrG1  | nProLeuTyr | ArgGlyGluL | euAsnGluHi  |            |      |
| TTTGGGACTC | CTGGGGCCAT  | ATATAAGAGC | AGAAGTTGAA | GATAATATCA  |            |      |
| sLeuGlyLeu | LeuGlyProt  | yrIleArgAl | aGluValGlu | AspAsnIleM  |            |      |
| TGGTAACTTT | CAGAAATCAG  | GCCTCTCGTC | CCTATTCTT  | CTATTCTAGC  |            |      |
| etValThrPh | eArgAsnGln  | AlaSerArgP | roTyrSerPh | eTyrSerSer  |            |      |
| CITATTCTT  | ATGAGGAAGA  | TCAGAGGCAA | GGAGCAGAAC | CTAGAAAAAA  |            |      |
| LeuIleSerT | yrGluGluAs  | pGlnArgGln | GlyAlaGluP | roArgLysAs  |            |      |
| CTTGTCAG   | CCTAATGAAA  | CCAAAACITA | CITTTGGAAA | GIGCAACATC  |            |      |
| nPheValLys | ProAsnGLUT  | hrLysThrTy | rPheTrpLys | ValGlnHisH  |            |      |
| ATATGGCACC | CACTAAAGAT  | GAGTTGACT  | GCAAAGCCTG | GCCTTATTTC  |            |      |
| isMetAlaPr | oThrLysAsp  | GluPheAspC | ysLysAlaTr | pAlaTyrPhe  |            |      |
| TCTGATGTG  | ACCTGGAAAA  | AGATGIGCAC | TCAGGCCTGA | TTGGACCCCT  |            |      |
| SerAspValA | spLeuGluLy  | sAspValHis | SerGlyLeuI | leGlyProLe  |            |      |
| TCTGGCTCTG | CACACTAACAA | CACTGAACCC | TGCTCATGGG | AGACAAGTGA  |            |      |
| uLeuValCys | HisThrAsnT  | hrLeuAsnPr | oAlaHisGly | ArgGlnValT  |            |      |
| CAGTACAGGA | ATTIGCTCTG  | TTTTTCACCA | TCTTGATGA  | GACCAAAAGC  |            |      |
| hrValGlnG1 | uPheAlaLeu  | PhePheThrI | lePheAspG1 | uThrLysSer  |            |      |
| TGGTACTTCA | CTGAAAATAT  | GGAAAGAAAC | TGCAGGGCTC | CCTGCAATAT  |            |      |
| TrpTyrPheT | hrGluAsnMe  | tGluArgAsn | CysArgAlaP | roCysAsnI1  |            |      |
| CCAGATGGAA | GATCCCACIT  | TTAAAGAGAA | TTATCGCTTC | CATGCAATCA  |            |      |
| eGlnMetGlu | AspProThrP  | heLysGluAs | nTyrArgPhe | HisAlaIleA  |            |      |
| ATGGCTACAT | AATGGATACA  | CTACCTGGCT | TAGTAATGGC | TCAGGATCAA  |            |      |
| snglyTyrI1 | eMetAspIhr  | LeuProGlyL | euValMetAl | aGlnAspGlnO |            |      |
| AGGATTGAT  | GGTATCTGCT  | CAGCATGGGC | AGCAATGAAA | ACATOC      |            |      |
| ArgIleArgT | rpTyrIleLe  | uSerMetGly | SerAsnGluA | snJ         | 3800       |      |
| TATTCATTTC | AGTGGACATG  | TGTTCACTGT | ACGAAAAAAA |             |            |      |
| rIleHisPhe | SerGlyHisV  | alPheThrVa | 1ArgLys    |             |            |      |
| AAATGGCACT | GTACAACTTC  | TATOCAGGTG | TTT        |             |            |      |
| ysMetAlaLe | uTyrAsnLeu  | TyrProGlyV | a          |             |            |      |

FIGURE 1 (cont.)

| 10                | 20                | 30                | 40                | 50                |      |
|-------------------|-------------------|-------------------|-------------------|-------------------|------|
| <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> |      |
| TTAOCATCCA        | AAGCTGGAAT        | TGGGGGGTG         | GAATGCCCCA        | TTGGGGAGCA        | 3850 |
| LeuProSerL        | ysAlaGlyI1        | eTrpArgVal        | GluCysLeuI        | leGlyGluHi        |      |
| TCTACATGCT        | GGGATGAGCA        | CACTTTCT          | GGTGTACAGC        | AATAAGTGTC        | 3900 |
| sLeuHisAla        | GlyMetSerT        | hrLeuPheLe        | uValTyrSer        | AsnLysCysG        |      |
| AGACTCCCCCT       | GGGAATGGCT        | TCTGGACACA        | TTAGAGATT         | TCAGATTACA        | 3950 |
| 1nThrProLe        | uGlyMetAla        | SerGlyHisI        | leArgAspPh        | eGlnIleThr        |      |
| GCTTCAGGAC        | AATATGGACA        | GTGGGCCCCA        | AAGCTGGCCA        | GACTTCATTA        | 4000 |
| AlaSerGlyG        | 1nTyrGlyG1        | nTrpAlaPro        | LysLeuAlaA        | ngLeuHisTy        |      |
| TTCGGGATCA        | ATCAATGCCT        | GGAGCACCAA        | GGAGGCCCTT        | TCTTGGATCA        | 4050 |
| rSerGlySer        | IleAsnAlaT        | rpSerThrLy        | sGluProPhe        | SerTrpIleL        |      |
| AGGIGGATCT        | GTTGGCACCA        | ATGATTATTC        | ACGGCATCAA        | GACCCAGGGT        | 4100 |
| ysValAspLe        | uLeuAlaPro        | MetIleIleH        | isGlyIleLy        | sThrGlnGly        |      |
| GCCCGTCAGA        | AGITCTCCAG        | CCTCTACATC        | TCTCAGTTTA        | TCATCATGTA        | 4150 |
| AlaArgGlnL        | ysPheSerSe        | rLeuTyrIle        | SerGlnPheI        | leIleMetTy        |      |
| TAGCTTGTAT        | GGGAAGAAGT        | GGCAGACTTA        | TCGAGGAAAT        | TOCACTGGAA        | 4200 |
| rSerLeuAsp        | GlyLysLysT        | rpGlnThrTy        | rArgGlyAsn        | SerThrGlyT        |      |
| CCTTAATGGT        | CTTCTTTGGC        | AATGTGGATT        | CATCTGGGAT        | AAAACACAAT        | 4250 |
| hrLeuMetVa        | 1PhePheGly        | AsnValAspS        | erSerGlyI1        | eLysHisAsn        |      |
| ATTTTTAACCC       | CTCCAATTAT        | TGCTCGATAC        | ATCCGGTTGC        | ACCCAACCTCA       | 4300 |
| IlePheAsnP        | roProIleI1        | eAlaArgTyr        | IleArgLeuH        | isProThrHi        |      |
| TTATAGCATT        | CGCAGCACTC        | TTCGCATGGA        | GTGATGGC          | TGTGATTAA         | 4350 |
| sTyrSerIle        | ArgSerThrL        | euArgMetG1        | uLeuMetGly        | CysAspLeuA        |      |
| ATAGTTGCAG        | CATGCCATTG        | CGAATGGAGA        | GTAAAGCAAT        | ATCAGATGCA        | 4400 |
| snSerCysSe        | rMetProLeu        | GlyMetGluS        | erLysAlaI1        | eSerAspAla        |      |
| CAGATTACTG        | CTTCATCTA         | CTTACCAAT         | ATGTTGCCA         | CCTGGCTCTCC       | 4450 |
| GlnIleThrA        | 1aSerSerTy        | rPheThrAsn        | MetPheAlaT        | hrTrpSerPr        |      |
| TTCAAAAGCT        | CGACTTCACC        | TOCAAGGGAG        | GAGTAATGCC        | TGGAGACCTC        | 4500 |
| oSerLysAla        | ArgLeuHisL        | euGlnGlyAr        | gSerAsnAla        | TrpArgProG        |      |
| AGGIGAATAA        | TCCAAAAGAG        | TGGCTGCAAG        | TGGACTTCCA        | GAAGACAATG        | 4550 |
| 1nValAsnAs        | nProLysGlu        | TrpLeuGlnV        | a1AspPheG1        | nLysThrMet        |      |
| AAAGTCACAG        | GAGTAACATAC       | TCAGGGAGTA        | AAATCTCTGC        | TTACCAGCAT        | 4600 |
| LysValThrG        | 1yValThrTh        | rGlnGlyVal        | LysSerLeuL        | euThrSerMe        |      |
| GTATGIGAAG        | GAGITCCCTCA       | TCTCCAGCAG        | TCAAGATGGC        | CATCAGTGGA        | 4650 |
| tTyrValLys        | GluPheLeuI        | leSerSerSe        | rGlnAspGly        | HisGlnTrpT        |      |
| CTCTCTTTT         | TCAGAATGGC        | AAAGTAAAGG        | TTTTTCAGGG        | AAATCAAGAC        | 4700 |
| hrLeuPhePh        | eGlnAsnGly        | LysValLysV        | a1PheGlnG1        | yAsnGlnAsp        |      |
| TCCTTCACAC        | CTGIGGGGAA        | CTCTCTAGAC        | CCACCGTTAC        | TGACTCGCTA        | 4750 |
| SerPheThrP        | roValValAs        | nSerLeuAsp        | ProProLeuL        | euThrArgTy        |      |

**FIGURE 1 (cont.)**

| 10                 | 20                | 30                | 40                 | 50                  |      |
|--------------------|-------------------|-------------------|--------------------|---------------------|------|
| <u>1234567890</u>  | <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u>  | <u>1234567890</u>   |      |
| CCTTCAATT          | CACCCCCAGA        | GTTGGGIGCA        | CCAGATTGCC         | CIGAGGATGG          | 4800 |
| rLeuArgIle         | HisProGlnS        | erTrpValHi        | sGlnIleAla         | LeuArgMetG          |      |
| AGGTCTGGG          | CTGCGAGGCA        | CAGGACCTCT        | ACTGACTCGA         | GCGAGTTCTT          | 4850 |
| luValLeuGl         | yCysGluAla        | GlnAspLeuT        | yr...              |                     |      |
| CTGAGGGAT          | CGGCAATAAA        | AAGACAGAAT        | AAAACGCACG         | GGTGTGGGT           | 4900 |
| CGTTTGTTCG         | GATCCAGATC        | TAGGAACCCC        | TAGTGTATGGA        | GTTGGCCACT          | 4950 |
| CCCTCTCTGC         | GCGCTCGCTC        | GCTCACTGAG        | GGGGGGGGGG         | CAAAGCCCCG          | 5000 |
| GCGTGGGAG          | ACCTTGGTC         | GGCGGCGCTC        | AGTGAGCGAG         | CGAGCGCGCA          | 5050 |
| GAGAGGGAGT         | GGCCAACCCC        | CCCCCCCCCC        | CCCTGCAGC          | CCAGCTGCAT          | 5100 |
| TAATGAATCG         | GCCAAACGCGC       | GGGGAGAGGC        | GGTTTGCATA         | TTGGGAGCTC          | 5150 |
| <u>TTCCGCTTCC</u>  | <u>TCGCTCACTG</u> | <u>ACTCGCTGCG</u> | <u>CTGGTACGTT</u>  | <u>CGGCTGCGC</u>    | 5200 |
| <u>GAGCGGTATC</u>  | <u>AGCTCACTCA</u> | <u>AAGGCGGTAA</u> | <u>TACGGTTATC</u>  | <u>CACAGAACATCA</u> | 5250 |
| <u>GGGGATAACG</u>  | <u>CAGGAAAGAA</u> | <u>CATGTGAGCA</u> | <u>AAAGGCCAGC</u>  | <u>AAAAGGCCAG</u>   | 5300 |
| <u>GAACCGTAAA</u>  | <u>AAGGCGCGT</u>  | <u>TGCTGGCGTT</u> | <u>TTTCCATAGG</u>  | <u>CTCGGCCCCC</u>   | 5350 |
| <u>CTGAACGAGCA</u> | <u>TCACAAAAAT</u> | <u>CGACGCTCAA</u> | <u>GTCAAGAGGTG</u> | <u>GCGAAACCG</u>    | 5400 |
| <u>ACAGGACTAT</u>  | <u>AAAGATACCA</u> | <u>GGGGTTCC</u>   | <u>CCTGGAAAGCT</u> | <u>CCCTCGTGC</u>    | 5450 |
| <u>CTCTCCTGTT</u>  | <u>CGAACCTGC</u>  | <u>CGCTTACCGG</u> | <u>ATACCTGTC</u>   | <u>GCCTTCTCC</u>    | 5500 |
| <u>CTTGGGAAG</u>   | <u>CGTGGCGCTT</u> | <u>TCTCAATGCT</u> | <u>CAAGCTGTAG</u>  | <u>GTATCTCAGT</u>   | 5550 |
| <u>TCGGTGTAGG</u>  | <u>TCGTTGCTC</u>  | <u>CAAGCTGGGC</u> | <u>TGTTGCGACG</u>  | <u>AACCCCCCGT</u>   | 5600 |
| <u>TCAGCCCGAC</u>  | <u>CGCTGCGCCT</u> | <u>TATCCGGTAA</u> | <u>CTATCGTC</u>    | <u>TGAGTCCAACC</u>  | 5650 |
| <u>CGGTAAGACA</u>  | <u>CGACTTATCG</u> | <u>CCACTGGCAG</u> | <u>CAGCCACTGG</u>  | <u>TAACAGGATT</u>   | 5700 |

FIGURE 1 (cont.)

| 10                | 20                | 30                | 40                | 50                |      |
|-------------------|-------------------|-------------------|-------------------|-------------------|------|
| <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> |      |
| <u>AGCAGAGOGA</u> | <u>GGTATGTAGG</u> | <u>CGGTGCTACA</u> | <u>GAGTCTTGA</u>  | <u>AGTGGTGGC</u>  | 5750 |
| <u>TAACTAOGGC</u> | <u>TACACTAGAA</u> | <u>GGACAGTATT</u> | <u>TGGTATCTGC</u> | <u>GCTCTGCTGA</u> | 5800 |
| <u>AGCCAGTTAC</u> | <u>CTTCGGAAAA</u> | <u>AGAGTGGTA</u>  | <u>GCTCTGATC</u>  | <u>CGGCAAACAA</u> | 5850 |
| <u>ACCACCGCTG</u> | <u>GTACGGTGG</u>  | <u>TTTTTTGTT</u>  | <u>TGCAAGCAGC</u> | <u>AGATTACGCG</u> | 5900 |
| <u>CAGAAAAAAA</u> | <u>GGATCTCAAG</u> | <u>AAGATCCTTT</u> | <u>GATCTTTCT</u>  | <u>ACGGGGTCTG</u> | 5950 |
| <u>ACGCTCAGTG</u> | <u>GAACGAAAAC</u> | <u>TCACGTTAAG</u> | <u>GGATTTGGT</u>  | <u>CATGAGATTA</u> | 6000 |
| <u>TCAAAAGGA</u>  | <u>TCTCACCTA</u>  | <u>GATCCTTTA</u>  | <u>AATTAAAAAT</u> | <u>GAAGTTTAA</u>  | 6050 |
| <u>ATCAATCTAA</u> | <u>AGTATATATG</u> | <u>AGTAAACTTG</u> | <u>GTCTGACAGT</u> | <u>TACCAATGCT</u> | 6100 |
|                   |                   |                   |                   | y1GelIreS         |      |
| TAATCAGTGA        | GGCACCTATC        | TCAGCGATCT        | GTCTATTTCG        | TTCATCCATA        | 6150 |
| ueL...siHo        | rPlaV...gr        | AueLreSgrA        | psAelIulGn        | sAteMprTue        |      |
| GTTGCCTGAC        | TCCCGCTCGT        | GTAGATAACT        | ACGATAACGGG       | AGGGCTTACC        | 6200 |
| In1GgrAlaV        | y1GgrAgrAr        | hTreSueL...       | reSlaVorP         | orPreSlaVt        |      |
| ATCTGGCCCC        | AGTGCTGCAA        | TGATAACCGCG       | AGACCCACGC        | TCACCGGCTC        | 6250 |
| eMn1GylGpr        | TsiHn1GueL        | reSlaValAu        | eLy1GlaVre        | SlaVorPulG        |      |
| CAGATTATAC        | AGCAATAAAC        | CAGCCAGCG         | GAAGGGCGA         | GOGCAGAAAGT       | 6300 |
| ueLnsAelIu        | eLueLueLy1        | GalAueLgrA        | ehPorPgrAa        | 1AsyCehPsi        |      |
| GGTCTCTGCAA       | CTTTATCCGC        | CTCCATCCAG        | TCTATTAAATT       | GTTGCGGGGA        | 6350 |
| HpsAn1GueL        | syLelIgrAg        | rAprTylGrh        | T.....nsA         | nsAylGorPu        |      |
| AGCTAGAGTA        | AGTAGTTCGC        | CAGTTAATAG        | TTTGCACAC         | GTGTTGCCA         | 6400 |
| eL...ueLue        | LryThsAalA        | ueL...ryTn        | sAalAsyCgr        | An1Gn1GprT        |      |
| TTGCTACAGG        | CATCGTGGTG        | TCACGCTCGT        | CGTTTGGTAT        | GGCTTCATTC        | 6450 |
| n1G...ueLs        | yCgrAorPrh        | TlaVreSrht        | rhTh1GryTo        | rPsyLteM..        |      |
| AGCTCGGGTT        | CCCAAACGATC       | AAGGCCAGTT        | ACATGATCCC        | CCATGTTGTC        | 6500 |
| .reSgrAnsA        | y1GlaVelIu        | eLalAueL...       | teMeliylG         | prTrhTrhTs        |      |
| CAAAAAAGCG        | GTTAGCTCT         | TCGGTCTCTC        | GATCGTGTTC        | AGAAGTAAGT        | 6550 |
| yCehPueLor        | P...reSgrA        | grApsAulGr        | eSgrAn1G...       | ehPryTrht         |      |
| TGGCGCGAGT        | GTTATCACTC        | ATGGTTATGG        | CAGCACTGCA        | TAATTCTCTT        | 6600 |
| orPgrAueLr        | hTel1laV..        | .orP...orP        | ueLLaValAr        | yThsAulG..        |      |
| ACTGTCATGC        | CATCCGTAAG        | ATGCTTTCT         | GTGACTGGTG        | AGTACTCAAC        | 6650 |
| .n1G...alA        | teMgrAueLe        | 1IreSsyLnl        | GreSnlGsiH        | rhTreSueLp        |      |

FIGURE 1 (cont.)

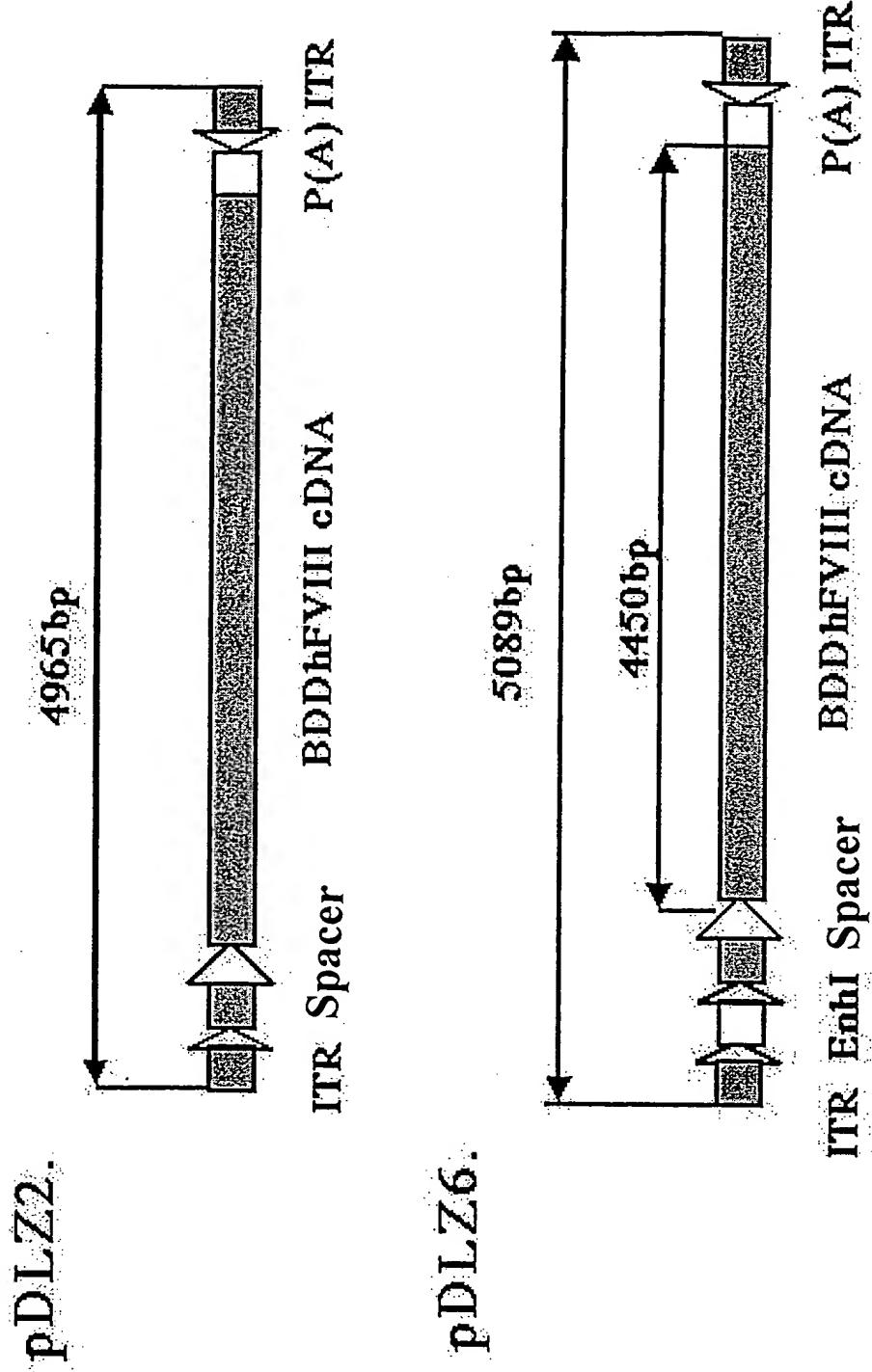
| 10                | 20                | 30                | 40                  | 50                |      |
|-------------------|-------------------|-------------------|---------------------|-------------------|------|
| <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u>   | <u>1234567890</u> |      |
| CAAGTCATTG        | TGAGAAATAGT       | GTATGCGGCG        | ACCGAGTTC           | TCTTGCGCGG        | 6700 |
| rTrhTteMgr        | AueLeIrhT         | ryTalAaIaI        | aVreSnsAre          | SsyLy1GorP        |      |
| CGTCAATAACG       | GGATAATACC        | GCGCACATA         | GCAGAACTTT          | AAAAGTGCTC        | 6750 |
| rhTueLlaVo        | rPryTryTgr        | AalAaVryT         | syCehPsyLu          | eLueLala..        |      |
| ATCATTGGAA        | AACGTTCTTC        | GGGGCGAAAA        | CTCTCAAGGA          | TCTTACCGCT        | 6800 |
| ....n1GehP        | 1aVnsAsyLo        | rPalAehPla        | VgrAueLreS          | grAlaValAr        |      |
| GTGAGATCC         | AGITCGATGT        | AACCCACTCG        | TGCACCCAAC          | TGATCTTCAG        | 6850 |
| hTreSelIpr        | ThsAreSrht        | 1aVprTulGs        | iHlaVprTre          | SelIsyLueL        |      |
| CATCCTTAC         | TTTCACCAAGC       | GTTCTGGGT         | GAGCAAAAC           | AGGAAGGCAA        | 6900 |
| teMsyL...s        | yL...prTgr        | AsyLn1GrhT        | ueLueLehPu          | eLehPalAeh        |      |
| AATGCGCAA         | AAAAGGGAAT        | AAGGGCGACA        | OGGAAATGTT          | GAATACTCAT        | 6950 |
| PsiHgrAueL        | ehPorPehPu        | eLorPreSla        | VreSelInsA          | ehPlaV...1        |      |
| ACTCTCCCTT        | TTTCAATATT        | ATTGAAGCAT        | TTATCAGGGT          | TATTGTCTCA        | 7000 |
| TGAGCGGATA        | CATAATTGAA        | TGTATTAGA         | AAAATAAAC           | AATAGGGTT         | 7050 |
| CGCGCACAT         | TTCCCCGAAA        | AGTGCACCT         | GACGTCTAAG          | AAACCATTAT        | 7100 |
| TATCATGACA        | TTAACCTATA        | AAAATAGGCG        | TATCACGAGG          | CCCTTCGTC         | 7150 |
| TCGCGCGTT         | CGGTGATGAC        | GGTGAAAACC        | TCTGACACAT          | GCAGCTCCCG        | 7200 |
| GAGACGGTCA        | CAGCTTGTCT        | GTAAGCGGAT        | GGGGGAGCA           | GACAAGCCCG        | 7250 |
| TCAGGGCGCG        | TCAGCGGGTG        | TGGCGGGTG         | TOGGGGCTGG          | CTTAACATATG       | 7300 |
| CGGCATCAGA        | GCAGATTGTA        | CTGAGAGTGC        | ACCATATGCG          | GTGTGAAATA        | 7350 |
| CGCACAGAT         | CGTAAGGAG         | AAAATACCGC        | <u>ATCAGGAAAT</u>   | <u>TGTAAACGTT</u> | 7400 |
| <u>AATATTGT</u>   | <u>TAAAATTGCG</u> | <u>GTTAAATT</u>   | <u>TTTGTTAAATCA</u> | <u>GCTCATTTT</u>  | 7450 |
| <u>TAACCAATAG</u> | <u>GGGAAATCG</u>  | <u>GCAAAATCC</u>  | <u>TTATAAAATCA</u>  | <u>AAAGAATAGA</u> | 7500 |
| <u>CGAGATAGG</u>  | <u>GTTGAGTGT</u>  | <u>GTTCCAGTTT</u> | <u>GGAAACAAGAG</u>  | <u>TCCACTATTA</u> | 7550 |
| <u>AAGAACGTGG</u> | <u>ACTCCAACGT</u> | <u>CAAAGGGCGA</u> | <u>AAAACCGTCT</u>   | <u>ATCAGGGCGA</u> | 7600 |

## ●FIGURE 1 (cont.)

| 10                 | 20                 | 30                 | 40                 | 50                 |      |
|--------------------|--------------------|--------------------|--------------------|--------------------|------|
| <u>1234567890</u>  | <u>1234567890</u>  | <u>1234567890</u>  | <u>1234567890</u>  | <u>1234567890</u>  |      |
| <u>TGGCCCCACTA</u> | <u>CGTGAACCAT</u>  | <u>CACCCCTAAC</u>  | <u>T AAGTTTTTG</u> | <u>GGGTGAGGT</u>   | 7650 |
| <u>GCGGTAAAGC</u>  | <u>ACTAAATCGG</u>  | <u>AACCCCTAAAG</u> | <u>GGAGCCCCCG</u>  | <u>ATTTAGAGCT</u>  | 7700 |
| <u>TGACGGGGAA</u>  | <u>AGCCGGCGAA</u>  | <u>CGTGGCGAGA</u>  | <u>AAGGAAGGGA</u>  | <u>AGAAAGCGAA</u>  | 7750 |
| <u>AGGAGGGGGC</u>  | <u>GCTAGGGCGC</u>  | <u>TGGCAAGTGT</u>  | <u>AGCGGTCAAG</u>  | <u>CTGCGCGTAA</u>  | 7800 |
| <u>CCACACACACC</u> | <u>CGCGCGCGCTT</u> | <u>AATGCGCGCG</u>  | <u>TACAGGGCGC</u>  | <u>GTCGCGCGCAT</u> | 7850 |
| <u>TOGCCATTCA</u>  | <u>GGCTAOGCAA</u>  | <u>CTGTTGGGAA</u>  | <u>GGCGGATCGG</u>  | <u>TGCGGGCGTC</u>  | 7900 |
| <u>TTCGCTATTAA</u> | <u>CGCGCAGCTGG</u> | <u>CTGCAGGGGG</u>  | <u>GGGGGGGGGG</u>  | <u>GGGT</u>        | 7944 |

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FIGURE 2



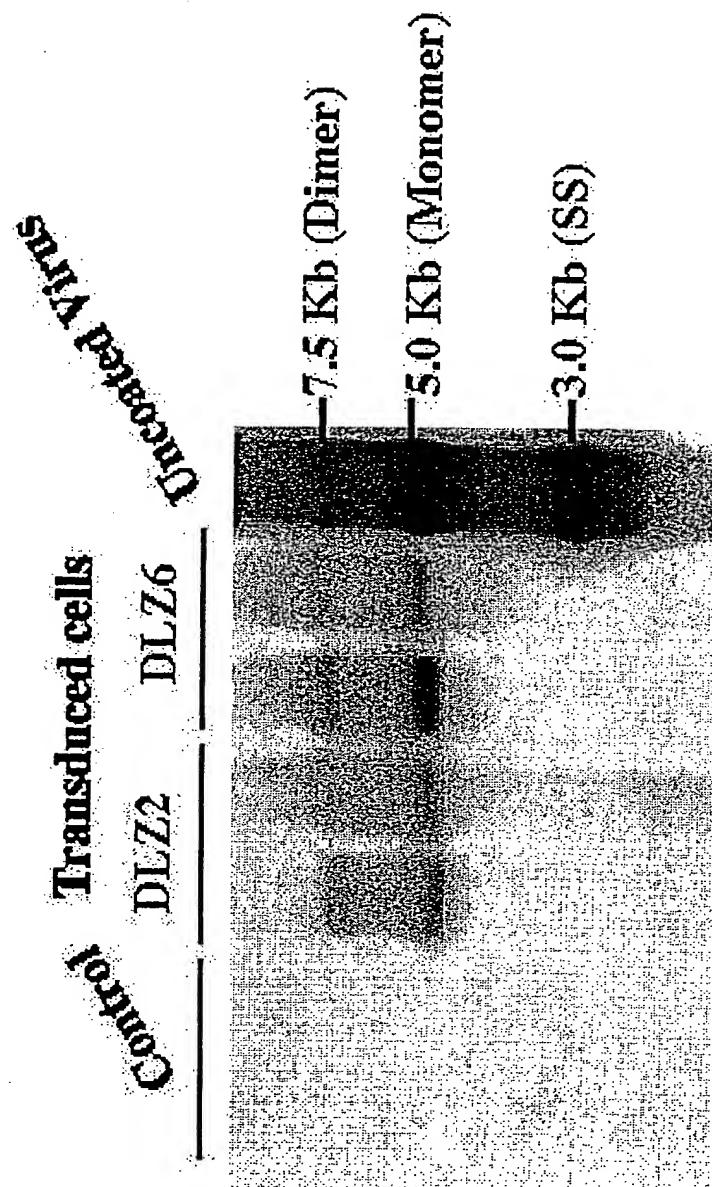


FIGURE 4-A

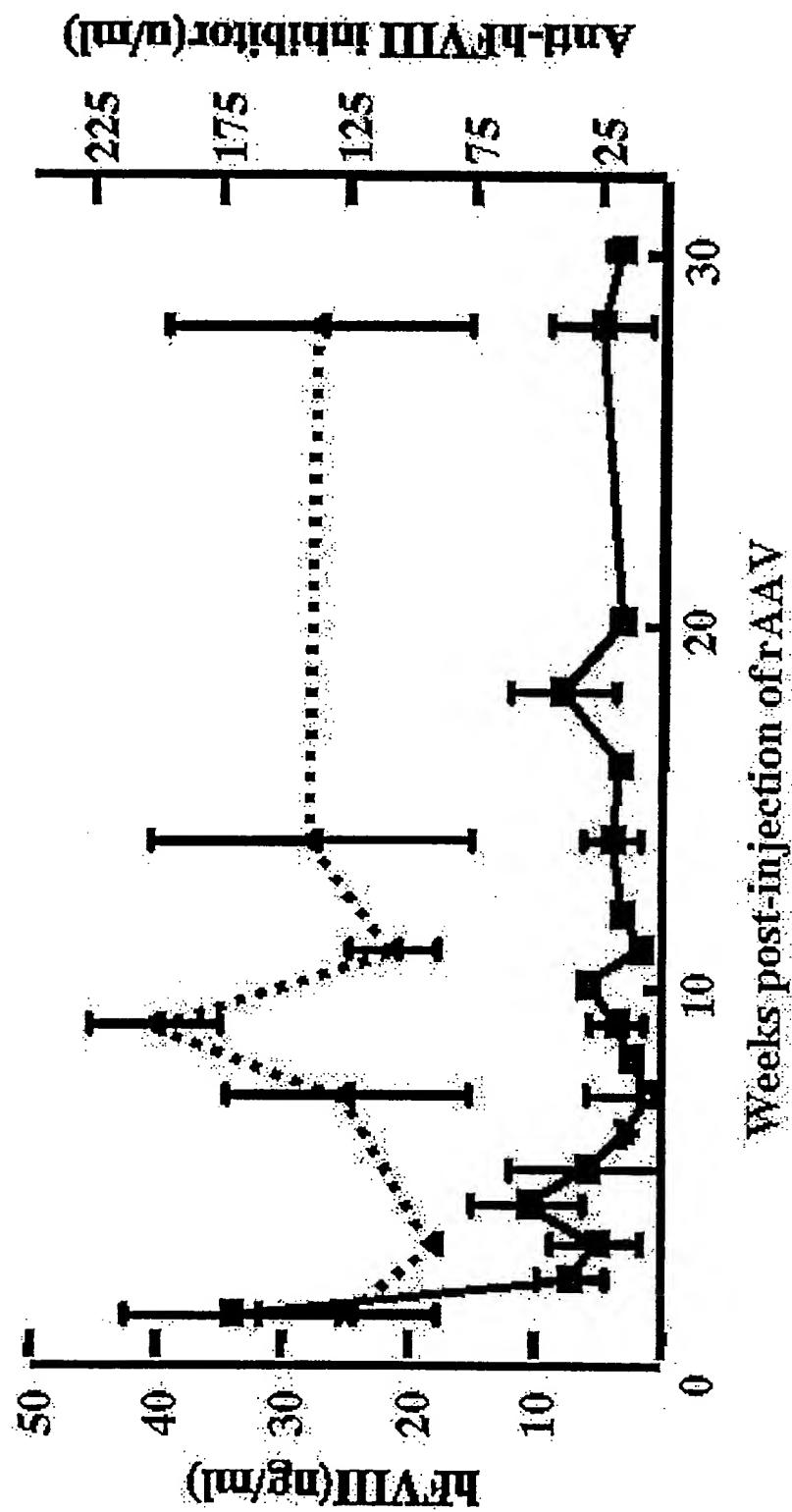


FIGURE 4-B

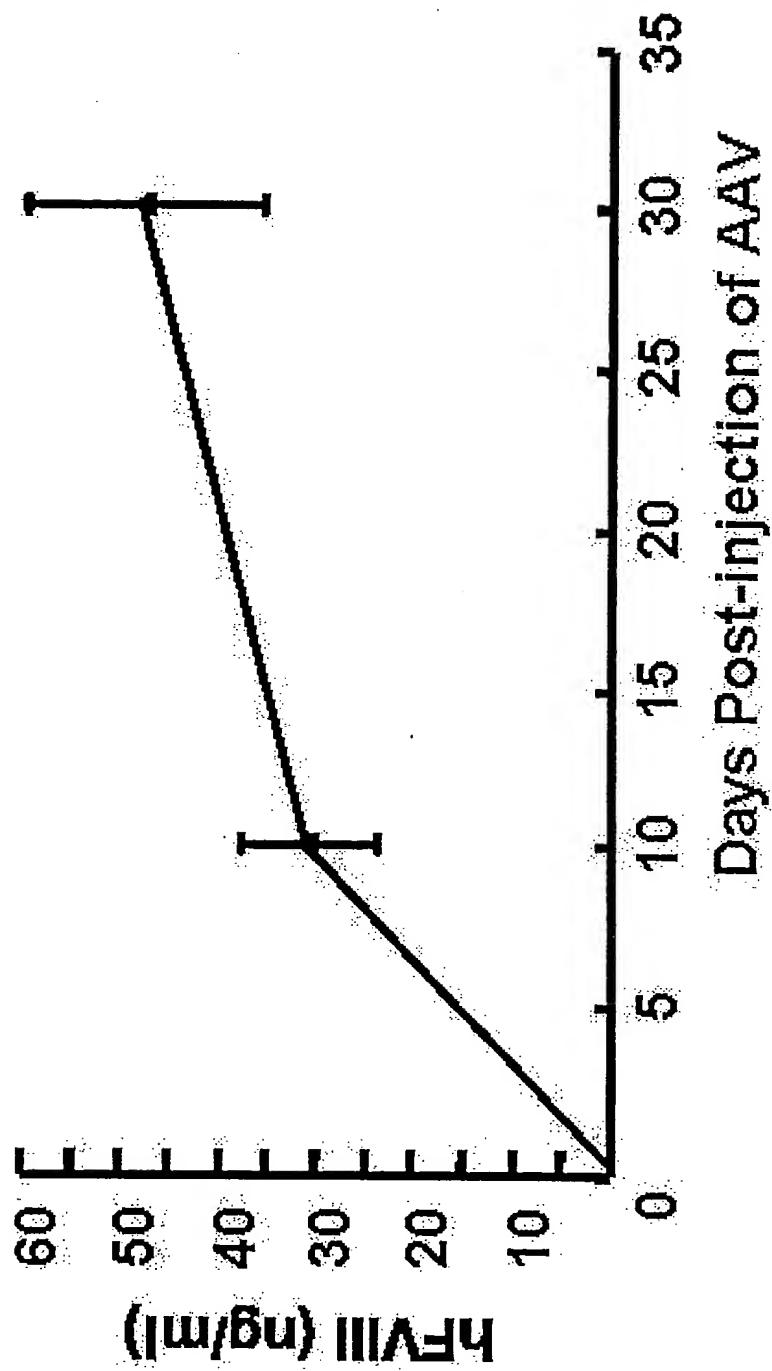


FIGURE 5-A

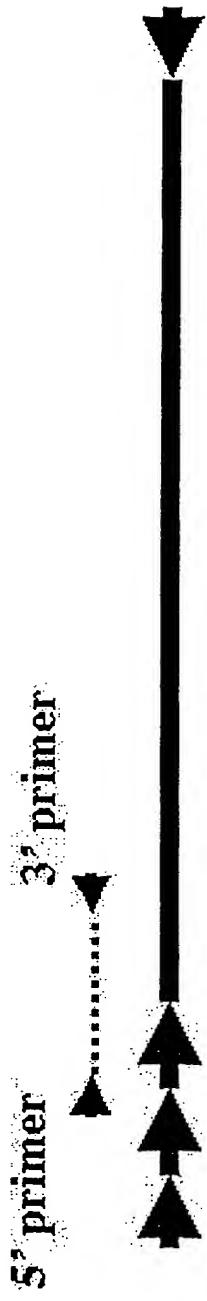


FIGURE 5-B

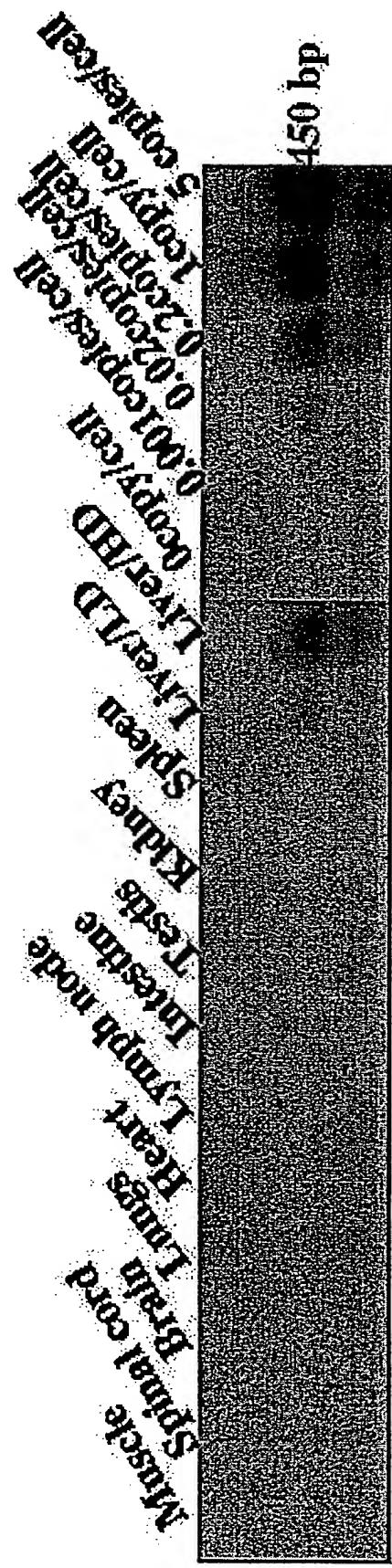


FIGURE 5-C

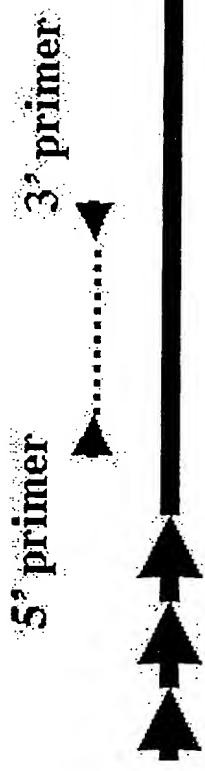
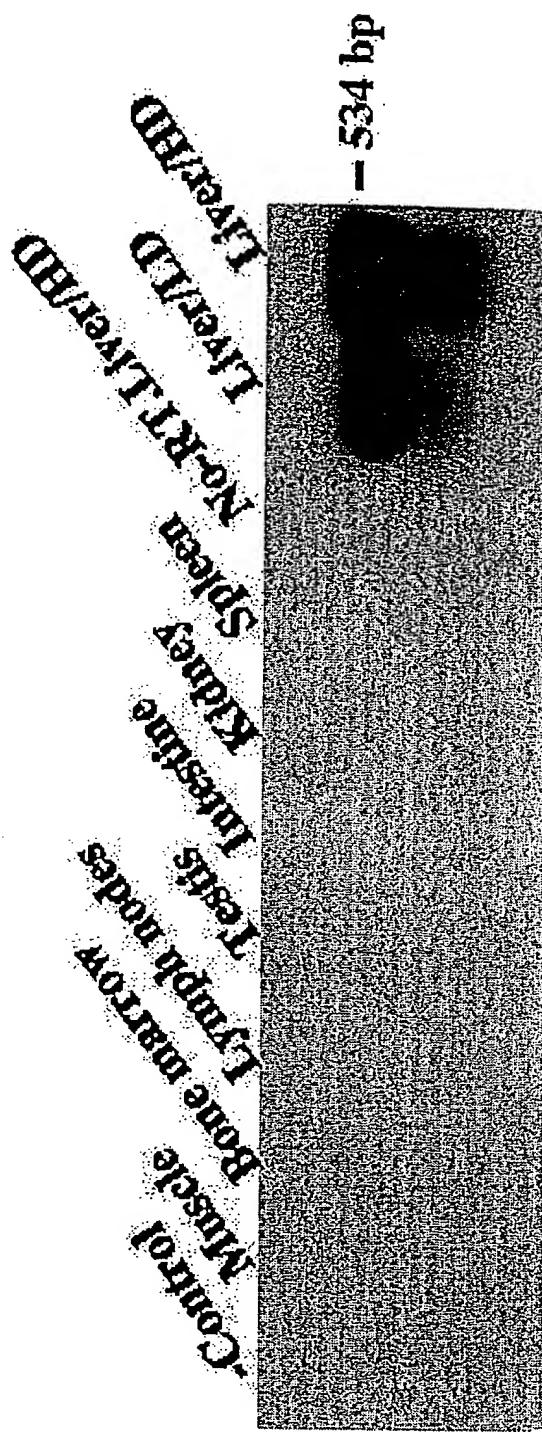


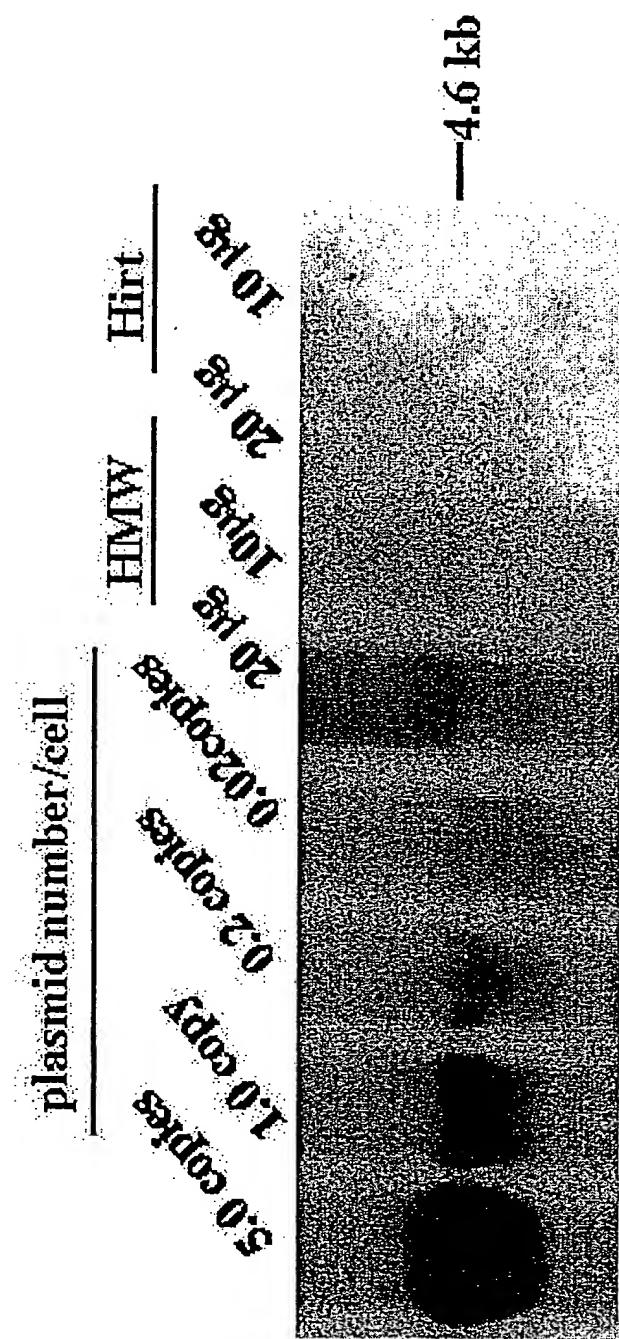
FIGURE 5-D



## FIGURE 5-E



FIGURE 5-F



# FIGURE 6

| 10                | 20                | 30                | 40                | 50                |     |
|-------------------|-------------------|-------------------|-------------------|-------------------|-----|
| <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> |     |
| TGGCCACTCC        | CTCTCTGCGC        | GCTGGCTCGC        | TCACTGAGGC        | CGGGCGACCA        | 50  |
| AAGGTGGGCC        | GAOGGCGGGG        | CTTTGGCGGG        | GCGGCGTCAG        | TGAGCGAGCG        | 100 |
| AGCGCGCAGA        | GAGGGAGTGG        | CCAACCTCCAT       | CACTAGGGGT        | TCTTCAGATC        | 150 |
| TCTTCTAAG         | TAAACAGTAC        | ATGAACCTTT        | ACCCCGITGC        | TGGCAACGG         | 200 |
| CCTGGCTCTG        | GCAGAAGTGT        | TGCTGACGCA        | ACCCCCACTG        | GCTGGGGCTT        | 250 |
| GGCGATAGGC        | CATCAGCGCA        | TGCGGATCTC        | AGTGTGGTT         | TGCAAGAGGA        | 300 |
| AGCAAAAAGC        | CTCTCCACCC        | AGGCCTGGAA        | TGTTTCCACC        | CAATGTCGAG        | 350 |
| CAGTGIGGTT        | TTGCAAGAGG        | AAGCAAAAAG        | CCCTCTCCACC       | CAGGCGCTGGA       | 400 |
| CTCGACCTCG        | AGAGTACTTC        | TAGAAATAACG       | AGCGATGCAA        | GTAGAGCTCT        | 450 |
|                   |                   |                   | MetGln            | ValGluLeuT        |     |
| ACACCTGCTG        | CCTTCTGTGC        | CTTTTGCCCT        | TCAGCCTTAG        | TGCCACCAGA        | 500 |
| yrThrCysCy        | sPheLeuCys        | LeuLeuProP        | heSerLeuSe        | rAlaThrArg        |     |
| AAATACTACC        | TCGGTGCAGT        | GGAACTGTCC        | TGGGACTATA        | TGCAAAGTGA        | 550 |
| LysTyrTyrL        | euGlyAlaVa        | 1GluLeuSer        | TrpAspTyrM        | etGlnSerAs        |     |
| CCTGCTCAGT        | GGCTGGCACG        | CGGATACAAG        | CTTTCTTCC         | AGGGTGCCAG        | 600 |
| pLeuLeuSer        | AlaLeuHisA        | 1aAspThrSe        | rPheSerSer        | ArgValProG        |     |
| GATCTTGGCC        | ACTCACCAOG        | TCAGTCACGT        | ACAGAAAGAC        | TGTGTTTGT         | 650 |
| lySerLeuPr        | oLeuThrThr        | SerValThrT        | yrArgLysTh        | rValPheVal        |     |
| GAGTTTACAG        | ATGACCTTT         | CAACATTGCC        | AAGCCCAGGC        | CACCGTGGAT        | 700 |
| GluPheThrA        | spAspLeuPh        | eAsnIleAla        | LysProArgP        | roProTrpMe        |     |
| GGGCCTGCTG        | GGTCTTACCA        | TOCAGGCTGA        | GGTTTATGAC        | ACAGGGTCA         | 750 |
| tGlyLeuLeu        | GlyProThrI        | 1eGlnAlaG1        | uValTyrAsp        | ThrValValI        |     |
| TGTCCTTAA         | GAACATGGCT        | TCTCATCTG         | TCAGCCTTCA        | CGCTGTTGGT        | 800 |
| 1eValLeuLy        | sAsnMetAla        | SerHisProV        | alSerLeuHi        | sAlaValGly        |     |
| GTATCCTATT        | GGAAAGCTTC        | TGAAGGIGCT        | GAGTATGAGG        | ATCAGACCAAG       | 850 |
| ValSerTyrT        | rpLysAlaSe        | rGluGlyAla        | GluTyrGluA        | spGlnThrSe        |     |
| CCAAAAGGAG        | AAGGAAGATG        | ATAATGTCAT        | TCCTGGTGA         | AGCCATAACCT       | 900 |
| rGlnLysGlu        | LysGluAspA        | spAsnValII        | eProGlyGlu        | SerHisThrT        |     |
| ATGTCTGGCA        | GGTCTTGTAAA       | GAGAATGCC         | CAATGGCCTC        | TGATCCACCA        | 950 |
| yrValTrpG1        | nValLeuLys        | GluAsnGlyP        | roMetAlaSe        | rAspProPro        |     |

● FIGURE 6 (cont.)

| 10                | 20                | 30                | 40                | 50                |      |
|-------------------|-------------------|-------------------|-------------------|-------------------|------|
| <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> |      |
| TGTCTCACCT        | ACTCATATT         | TTCACACGTG        | GACCTGGTGA        | AAGACCTGAA        | 1000 |
| CysLeuThrT        | yrSerTyrPh        | eSerHisVal        | AspLeuValL        | ysAspLeuAs        |      |
| TTCAGGCTC         | ATTGGAGGCC        | TGCTGGTTTG        | CAAAGAAGGG        | AGTCTGGCCA        | 1050 |
| nSerGlyLeu        | IleGlyAlaL        | euLeuValCy        | sLysGluGly        | SerLeuAlaL        |      |
| AAGAAAGGAC        | ACAGACCTTG        | CAGGAATTIG        | TCCTACTTT         | TGCTGTATT         | 1100 |
| ysGluArgTh        | rGlnThrLeu        | GlnGluPheV        | alLeuLeuPh        | eAlaValPhe        |      |
| GATGAAGGGA        | AAAGTTGGCA        | CTCAGAAACA        | AATGCGCTT         | TGACACAGGC        | 1150 |
| AspGluGlyL        | ysSerTrpHi        | sSerGluThr        | AsnAlaSerL        | euThrGlnAl        |      |
| TGAGGOCAG         | CATGAGCTGC        | ACAOCCATCAA       | TGGCTATGTA        | AACAGGTCTC        | 1200 |
| aGluAlaGln        | HisGluLeuH        | isThrIleAs        | nGlyTyrVal        | AsnArgSerL        |      |
| TGCGAGGCT         | TACTGIGGTT        | CACAAGAGAT        | CAGTCTATTG        | GCATGIGATT        | 1250 |
| euProGlyLe        | uThrValCys        | HisLysArgS        | erValTyrTr        | pHisValIle        |      |
| GGAATGGGCA        | CCACCCCCGA        | AGTGCCTCA         | ATTTTCTCG         | AAGGTACAC         | 1300 |
| GlyMetGlyT        | hrThrProG1        | uValHisSer        | IlePheLeuG        | luGlyHisTh        |      |
| ATTCTCTGIG        | AGGAACCACC        | GCCAGGCTC         | CTTGGAGATC        | TCACCAATT         | 1350 |
| rPheLeuVal        | ArgAsnHisA        | rgGlnAlaSe        | rLeuGluIle        | SerProIleT        |      |
| CITTCCTTAC        | TGCTCAGACA        | TTCCCTGATGG       | ACCTGGCCA         | GTTCCTACTG        | 1400 |
| hrPheLeuTh        | rAlaGlnThr        | PheLeuMetA        | spLeuGlyG1        | nPheLeuLeu        |      |
| TTTGTCTATA        | TCCCTTCCC         | TCAACATGAT        | GGTATGGAAG        | CTTATGTCAA        | 1450 |
| PheCysHisI        | leProSerHi        | sGlnHisAsp        | GlyMetGluA        | laTyrValLy        |      |
| AGTAGATAGC        | TGCCCCAGAGG       | AAACCCAGCT        | GCGCATGAAA        | AATAATGAAG        | 1500 |
| sValAspSer        | CysProGluG        | luProGlnLe        | uArgMetLys        | AsnAsnGluA        |      |
| ATAAAAGATTA       | TGATGATGGT        | CTTTATGATT        | CTGACATGGA        | CGTAGITAGC        | 1550 |
| spLysAspTy        | rAspAspGly        | LeuTyrAspS        | erAspMetAs        | pValValSer        |      |
| TTTGTGACG         | ACAGCTCTTC        | TCCCTTATC         | CAAATCCGCT        | CAGTTGCCAA        | 1600 |
| PheAspAspA        | spSerSerSe        | rProPheIle        | GlnIleArgS        | erValAlaLy        |      |
| GAAGCATCCT        | AAAACCTGGG        | TCCACTATAT        | TGCTGCTGAG        | GAGGAGGACT        | 1650 |
| sLysHisPro        | LysThrTrpV        | alHisTyrIl        | eAlaAlaGlu        | GluGluAspT        |      |
| GGGACTATGC        | TOCCCTCAGGC       | CCCACCCCCA        | ATGATAGAAG        | TCATAAAAAT        | 1700 |
| rpAspTyrAl        | aProSerGly        | ProThrProA        | snAspArgSe        | rHisLysAsn        |      |
| CTGTATTGTA        | ACAATGGTCC        | TCAGGGATT         | GGTAAGAAGT        | ACAAAAAAAGT       | 1750 |
| LeuTyrLeuA        | snAsnGlyPr        | oGlnArgIle        | GlyLysLysT        | yrLysLysVa        |      |
| CCGATTGIG         | GCATACACAG        | ATGAGACATT        | TAAGACTCGT        | GAAGCTATTC        | 1800 |
| 1ArgPheVal        | AlaTyrThrA        | spGluThrPh        | eLysThrArg        | GluAlaIleG        |      |
| AGTATGAATC        | AGGAATCTG         | GGACCTTAC         | TTTATGGAGA        | AGTTGGAGAC        | 1850 |
| lnTyrGluSe        | rGlyIleLeu        | GlyProLeuL        | euTyrGlyG1        | uValGlyAsp        |      |
| ACACTGCTGA        | TTATATTAA         | GAATCAAGGCC       | AGCGGGCCAT        | ATAAACATCTA       |      |
| ThrLeuLeuI        | leIlePheLy        | sAsnGlnAla        | SerArgProT        | yrAsnIleTy        | 1900 |

FIGURE 6 (cont.)

| 10   | 20         | 30         | 40         | 50         |
|--|------------|------------|------------|------------|
| 1234567890   | 1234567890 | 1234567890 | 1234567890 | 1234567890 |
| CCCTCATGGG ATCAAATTATG TCACTCTCT GCACACAGGG AGATTGCCAA   |            |            |            | 1950       |
| rProHisGly IleAsnTyrV alThrProLe uHisThrGly ArgLeuProL   |            |            |            |            |
| AAGGTGTGAA ACATTGAAA GATATGCCAA TTCTGCGGGG AGAGATATTC    |            |            |            | 2000       |
| ysGlyValLy sHisLeuLys AspMetProI 1eLeuProG1 yGluIlePhe   |            |            |            |            |
| AAGTATAAAT GGACAGTGAC CGTAGAAGAT GGACCAACTA AATCAGATCC   |            |            |            | 2050       |
| LysTyrLysT rpThrValTh rValGluAsp GlyProThrL ysSerAspPr   |            |            |            |            |
| TCGGTGCCTG ACCCGATATT ACTCAAGCTT CATTAATCTG GAGAGAGATC   |            |            |            | 2100       |
| oArgCysLeu ThrArgTyrT yrSerSerPh eIleAsnLeu GluArgAspL   |            |            |            |            |
| TAGCTTCAGG ACTCATTGGC CCTCTCTCA TCTGCTACAA AGAACCTGTA    |            |            |            | 2150       |
| euAlaSerG1 yLeuIleGly ProLeuLeuI 1eCysTyrLy sGluSerVal   |            |            |            |            |
| GATCAAAGAG GAAACCAGAT GATGTCAGAC AAGAGAAATG TCATCCTGTT   |            |            |            | 2200       |
| AspGlnArgG lyAsnGlnMe tMetSerAsp LysArgAsnV alIleLeuPh   |            |            |            |            |
| TCTGTATTT GATGAGAACG GAAGCTGGTA CCTCACAGAG AATATGCAGC    |            |            |            | 2250       |
| eSerValPhe AspGluAsnA rgSerTrpTy rLeuThrGlu AsnMetGlnA   |            |            |            |            |
| GCTTCCTCCC CAATGCAGAT GTAGTGCAGC CCCATGACCC AGAGTCCAA    |            |            |            | 2300       |
| rgPheLeuPr oAsnAlaAsp ValValGlnP roHisAspPr oGluPheGln   |            |            |            |            |
| CCTCTAAACA TCATGCACAG CATCAATGGC TATGTTTTG ACAACTTGCA    |            |            |            | 2350       |
| LeuSerAsnI 1eMetHisSe rIleAsnGly TyrValPheA spAsnLeuG1   |            |            |            |            |
| GCTGTCAGTT TGTTTGCATG AGGIGGGGTA CTGGTACATT CTAAGTGTG    |            |            |            | 2400       |
| nLeuSerVal CysLeuHisG luValAlaTy rTrpTyrIle LeuSerValG   |            |            |            |            |
| GAGCACAAAC TGACTTCCTG TCTGCTCTCT TCTCTGGATA TACCTCAAA    |            |            |            | 2450       |
| lyAlaGlnIh rAspPheLeu SerValPheP heSerGlyTy rThrPheLys   |            |            |            |            |
| CACAAAATGG TCTATGAAGA CACACTTACC CCTCTOCCAT TCTCAGGAGA   |            |            |            | 2500       |
| HisLysMetV alTyrGluAs pThrLeuThr LeuPheProP heSerGlyG1   |            |            |            |            |
| AACTGCTTC ATGCAATGG AAAACCCAGG TCTGIGGGIT CTGGGGTGCC     |            |            |            | 2550       |
| uThrValPhe MetSerMetG luAsnProG1 yLeuTrpVal LeuGlyCysH   |            |            |            |            |
| ACAACTCAGA CTTCGGAAC AGAGGCATGA CAGCCCTACT GAAGGTTTCT    |            |            |            | 2600       |
| isAsnSerAs pPheArgAsn ArgGlyMetT hrAlaLeuLe uLysValSer   |            |            |            |            |
| AGTTGTAACA GGACATTGA TGATTATTAT GAGGACACAT ACGAAGATAT    |            |            |            | 2650       |
| SerCysAsnA rgAsnIleAs pAspTyrTyr GluAspThrT yrGluAspI1   |            |            |            |            |
| TCCAACTCCC CTGCTAAATG AAAACAATGT AATTAAACCT AGAACCTCT    |            |            |            | 2700       |
| eProThrPro LeuLeuAsnG luAsnAsnVa 1IleLysPro ArgSerPheS   |            |            |            |            |
| CCCGAGAATTC AAGGCACCCCT AGCACTAAGG AAAAGCAATT GAAAATGAAG |            |            |            | 2750       |
| erGlnAsnSe rArgHisPro SerThrLysG luLysGlnLe uLysMetLys   |            |            |            |            |
| AGAGAAGATT TTGACATCTA CGGGGACTAT GAAAATCAGG GCCTCCGCAG   |            |            |            | 2800       |
| ArgGluAspP heAspIleTy rGlyAspTyr GluAsnGlnG lyLeuArgSe   |            |            |            |            |
| CTTICAAAG AAAACACGAC ACTATTCAT TGCTGCACTG GAGCGTCTCT     |            |            |            | 2850       |
| rPheGlnLys LysThrArgH isTyrPheI1 eAlaAlaVal GluArqLeuT   |            |            |            |            |

FIGURE 6 (cont.)

| 10                | 20                | 30                | 40                | 50                |      |
|-------------------|-------------------|-------------------|-------------------|-------------------|------|
| <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> |      |
| GGGATTATGG        | GATGAGTAGA        | TCTCCOCATA        | TACTAAGAAA        | CAGGGCTCAA        | 2900 |
| rpAspTyrGl        | yMetSerArg        | SerProHisI        | 1eLeuArgAs        | nArgAlaGln        |      |
| AGTGGGGATG        | TOCAGCAGTT        | CAAGAAAGGTG       | GTTTCCAGG         | AATTTACTGA        | 2950 |
| SerGlyAspV        | alGlnGlnPh        | eLysLysVal        | ValPheGlnG        | luPheThrAs        |      |
| TGGATCCTTT        | ACTCAGCCCT        | TATACCGTGG        | AGAACTGAAT        | GAACACTTGG        | 3000 |
| pGlySerPhe        | ThrGlnProL        | euTyrArgGl        | yGluLeuAsn        | GluHisLeuG        |      |
| GACTCTGGG         | GCCATATATA        | AGAGCAGAAG        | TTGAAGACAA        | TATCGTGGTA        | 3050 |
| 1yLeuLeuGl        | yProTyrIle        | ArgAlaGluV        | alGluAspAs        | nIleValVal        |      |
| ACTTCAAAA         | ACCAGGCTC         | TCGTCCTAC         | TCCTCTATT         | CTAGCTTAT         | 3100 |
| ThrPheLysA        | snGlnAlaSe        | rArgProTyr        | SerPheTyrS        | erSerLeuI1        |      |
| TTCTTATGAC        | GAAGATGAGG        | GACAAGGAGC        | AGAACCTAGA        | AGAAAGTTG         | 3150 |
| eSerTyrAsp        | GluAspGluG        | 1yGlnGlyAl        | aGluProArg        | ArgLysPheV        |      |
| TCAACCTAA         | TGAAACAAA         | ATTTACITTT        | GGAAAGTGCA        | GCATCATATG        | 3200 |
| alAsnProAs        | nGluThrLys        | IleTyrPheT        | rpLysValGl        | nHisHisMet        |      |
| GCACCCACTA        | AAGATGAGIT        | TGACTGCAA         | GCCTGGGCTT        | ATTTTCTGA         | 3250 |
| AlaProThrL        | ysAspGluPh        | eAspCysLys        | AlaTrpAlaT        | yrPheSerAs        |      |
| TGTTGATTG         | GAGAAAGATG        | TGCACTCAGG        | CTTGATTGGA        | CCCTTCTGA         | 3300 |
| pValAspLeu        | GluLysAspV        | alHisSerGl        | yLeuIleGly        | ProLeuLeuI        |      |
| TCTGCGCGAG        | TAACACACTG        | AACCTGCTC         | ATGGGAGACA        | AGTGACAGTG        | 3350 |
| 1eCysArgSe        | rAsnThrLeu        | AsnProAlaH        | isGlyArgGl        | nValThrVal        |      |
| CAGGAGTTG         | CCCTGGTTT         | CACTATATT         | GATGAGACTA        | AGAGCTGGTA        | 3400 |
| GlnGluPheA        | 1aLeuValPh        | eThrIlePhe        | AspGluThrL        | ysSerTrpTy        |      |
| CTTCACTGAA        | AACCTGGAAA        | GGAACTGTAG        | AGCTCCCTGC        | AATGTCCAGA        | 3450 |
| rPheThrGlu        | AsnLeuGluA        | rgAsnCysAr        | gAlaProCys        | AsnValGlnL        |      |
| AGGAGGACCC        | TACTCTAAA         | AAAAACTTCC        | GCTTCCATGC        | AATCAACGGC        | 3500 |
| ysGluAspPr        | oThrLeuLys        | GluAsnPheA        | rgPheHisAl        | aIleAsnGly        |      |
| TATGTGAAGG        | ATACACTOCC        | TGGCTTAGTA        | ATGGCTCAGG        | ATCAAAAGGT        | 3550 |
| TyrValLysA        | spThrLeuPr        | oGlyLeuVal        | MetAlaGlnA        | spGlnLysVa        |      |
| TCGATGGTAT        | CTGCTCAGCA        | TGGGCAGCAA        | CGAAAACATT        | CATTCCATT         | 3600 |
| 1ArgTrpTyr        | LeuLeuSerM        | etGlySerAs        | nGluAsnIle        | HisSerIleH        |      |
| ACTTCAGTGG        | ACATGTGTTC        | ACTGTACCGA        | AAAAAGAGGA        | ATATAAAATG        | 3650 |
| isPheSerGl        | yHisValPhe        | ThrValArgL        | ysLysGluGl        | uTyrLysMet        |      |
| GCAGTCTACA        | ACCTCTATCC        | AGGTGTTTT         | GAGACTGTGG        | AAATGCTAAC        | 3700 |
| AlaValTyrA        | snLeuTyrPr        | oGlyValPhe        | GluThrValG        | luMetLeuPr        |      |
| ATCCCCAAGTT       | GGAATCTGGC        | GGATAGAATG        | CCTTATCGGC        | GAGCACCTGC        | 3750 |
| oSerGlnVal        | GlyIleTrpA        | rgIleGluCy        | sLeuIleGly        | GluHisLeuG        |      |
| AAGCCGGGAT        | GAGCACTCTG        | TTTCTGGTGT        | ACAGCAAGAA        | GIGTCAGACT        | 3800 |
| lnAlaGlyMe        | tSerThrLeu        | PheLeuValT        | yrSerLysLy        | sCysGlnThr        |      |

FIGURE 6 (cont.)

| 10         | 20          | 30          | 40          | 50          |      |
|------------|-------------|-------------|-------------|-------------|------|
| 1234567890 | 1234567890  | 1234567890  | 1234567890  | 1234567890  |      |
| CCACTGGGGA | TGGCTTCGG   | ACACATTAGA  | GATTTTCAGA  | TTACAGCTTC  | 3850 |
| ProLeuGlyM | etAlaSerG1  | yHisIleArg  | AspPheGlnI  | leThrAlaSe  |      |
| AGGACAATAT | GGACAGTGGG  | CCCAAAGCT   | GGCCAGACTT  | CATTATTCG   | 3900 |
| rGlyGlnTyr | GlyGlnTrpA  | laProLysLe  | uAlaArgLeu  | HisTyrSerG  |      |
| GATCAATCAA | TGCTTGGAGC  | ACCAAGGATC  | CCCTTTCCTG  | GATCAAGGIG  | 3950 |
| lySerIleAs | nAlaTrpSer  | ThrLysAspP  | roPheSerTr  | pIleLysVal  |      |
| GATCTCTTGG | CACCGATGAT  | TATTCACGGC  | ATCATGACCC  | AGGGGGCCCG  | 4000 |
| AspLeuLeuA | laProMetI1  | eIleHisGly  | IleMetThrG  | lnGlyAlaAr  |      |
| CCAGAAGTTC | TOCAGCCCT   | ACGTGCTICA  | GTTTATCATC  | ATGTACAGTC  | 4050 |
| gGlnLysPhe | SerSerLeuT  | yrValSerG1  | nPheIleIle  | MetTyrSerL  |      |
| TGGATGGCAA | CAAGTGGCAC  | AGTTACCGAG  | GGAAATTCCAC | GGGGACCTTA  | 4100 |
| euAspGlyAs | nLysTrpHis  | SerTyrArgG  | lyAsnSerTh  | rGlyThrLeu  |      |
| ATGGCTCTCT | TIGGCAACGT  | GGATTACATCT | GGGATCAAAC  | ACAATATTT   | 4150 |
| MetValPheP | heGlyAsnVa  | 1AspSerSer  | GlyIleLysH  | isAsnIlePh  |      |
| TAACOCTCOG | ATTATTGCTC  | AGTACATCCG  | TTTGCACCCA  | ACCCATTACA  | 4200 |
| eAsnProPro | IleIleAlaG  | lnTyrIleAr  | gLeuHisPro  | ThrHisTyrS  |      |
| GCATCOGCAG | CACTCTTCGC  | ATGGAGCTCT  | TGGGCTGTGA  | CTTCAACAGT  | 4250 |
| erIleArgSe | rThrLeuArg  | MetGluLeuL  | euGlyCysAs  | pPheAsnSer  |      |
| TGCAGCATGC | CGCTGGGAT   | GGAGAGTAAA  | GCAATATCAG  | ATGCTCAGAT  | 4300 |
| CysSerMetP | roLeuGlyMe  | tGluSerLys  | AlaIleSerA  | spAlaGlnI1  |      |
| CACTGCTCTG | TCCTACCTAA  | GCAGTATGCT  | TGCCACTTGG  | TCTCCTTCCC  | 4350 |
| eThrAlaSer | SerTyrLeuS  | erSerMetLe  | uAlaThrTrp  | SerProSerG  |      |
| AAGCCCGGCT | GCACCTGGCAG | GGCAGGACTA  | ATGCCTGGAG  | ACCTCAGGCA  | 4400 |
| lnAlaArgLe | uHisLeuGln  | GlyArgThrA  | snAlaTrpAr  | gProGlnAla  |      |
| AATAACCAA  | AAGAGTGGCT  | GCAAGTGGAC  | TTCGGAAAGA  | CCATGAAAGT  | 4450 |
| AsnAsnProL | ysGluTrpLe  | uGlnValAsp  | PheArgLysT  | hrMetLysVa  |      |
| CACAGGAATA | ACCACCCAGG  | GGGTGAAATC  | TCTCCTCATC  | AGCATGTATG  | 4500 |
| lThrGlyIle | ThrThrGlnG  | lyValLysSe  | rLeuLeuIle  | SerMetTyrV  |      |
| TGAAGGAGTT | CCTCATCTCC  | AGTAGTCAAG  | ATGGCCATAA  | CTGGACTCTG  | 4550 |
| alLysGluPh | eLeuIleSer  | SerSerGlnA  | spGlyHisAs  | nTrpThrLeu  |      |
| TTTCTTCAGA | ATGGCAAAGT  | CAAGGTCTTC  | CAGGGAAACC  | GGGACTCTTC  | 4600 |
| PheLeuGlnA | snGlyLysVa  | 1LysValPhe  | GlnGlyAsnA  | rgAspSerSe  |      |
| CAACGCTGTG | CGGAACCGTC  | TGAAACCCCC  | GCTGGTGGCT  | CGCTACGTGTC | 4650 |
| rThrProVal | ArgAsnArgL  | euGluProPr  | oLeuValAla  | ArgTyrValA  |      |
| GCCTGCACCC | GCAGAGCTGG  | GCGCACCA    | TOGOCCTGAG  | GCTGGAGGTC  | 4700 |
| rgLeuHisPr | oGlnSerTrp  | AlaHisHisI  | leAlaLeuAr  | gLeuGluVal  |      |
| CTGGGCTGCG | ACACCCAGCA  | GCCCGCTGA   | CCGGCGCTC   | TGCGGCCCCG  | 4750 |
| LeuGlyCysA | spThrGlnG1  | nProAla...  |             |             |      |

## FIGURE 6 (cont.)

| 10                | 20                 | 30                 | 40                 | 50                |      |
|-------------------|--------------------|--------------------|--------------------|-------------------|------|
| <u>1234567890</u> | <u>1234567890</u>  | <u>1234567890</u>  | <u>1234567890</u>  | <u>1234567890</u> |      |
| TCTCCCTGCT        | CTCCTGCCC          | TGTCCCCGCG         | GCTCCCATC          | AAGCTTATCG        | 4800 |
| ATACCGTCGA        | GCGAGTTCTT         | CTGAGGGAT          | CGGCAATAAA         | AAGACAGAAT        | 4850 |
| AAAACGCAAG        | GGTGTGGGT          | CGTTTGTTCG         | GATCCAGATC         | TAGGAACCCC        | 4900 |
| TAGTGATGGA        | GTGGCCACT          | CCCTCTCTGC         | GCGCTGGCTC         | GCTCACTGAG        | 4950 |
| GCGGCGGGG         | CAAAGCCCGG         | GCGTGGGCG          | ACCTTGGTC          | GCGGGCGTC         | 5000 |
| AGTGAGGAG         | CGAGGCGGCA         | GAGAGGGAGT         | GGCCAACCCC         | CCCCCCCCCCC       | 5050 |
| CCCTGCAGC         | CCAGCTGCAT         | TAATGAATCG         | GCGAACGGCG         | GGGGAGAGGC        | 5100 |
| GGTTTGGTA         | TTGGGGCTC          | <u>TTCCGCTTCC</u>  | <u>TGGCTCACTG</u>  | <u>ACTGGCTGCG</u> | 5150 |
| <u>CTCGGTGTT</u>  | <u>CGGCTGGGC</u>   | <u>GAGCGGTATC</u>  | <u>AGCTCACTCA</u>  | <u>AAGGCGGTAA</u> | 5200 |
| <u>TACGGTTATC</u> | <u>CACAGAATCA</u>  | <u>GGGGATAACG</u>  | <u>CAGGAAAGAA</u>  | <u>CATGTGAGCA</u> | 5250 |
| <u>AAAGGCCAGC</u> | <u>AAAAGGCCAG</u>  | <u>GAACCGTAAA</u>  | <u>AAGGCCGCGT</u>  | <u>TGCTGGCGTT</u> | 5300 |
| <u>TTTCCATAGG</u> | <u>CTCCGGCCCCC</u> | <u>CTGACCGAGCA</u> | <u>TCACAAAAAT</u>  | <u>CGACGCTCAA</u> | 5350 |
| <u>GTCAGAGGTG</u> | <u>GCGAAACCGG</u>  | <u>ACAGGACTAT</u>  | <u>AAAGATAACCA</u> | <u>GGCGTTTCCC</u> | 5400 |
| <u>CCTGGAAGCT</u> | <u>CCCTCGTGGG</u>  | <u>CTCTCCGTT</u>   | <u>CGACCCCTGC</u>  | <u>CGCTTACCGG</u> | 5450 |
| <u>ATACCTGTC</u>  | <u>GCCTTCTCC</u>   | <u>CTTGGGAAG</u>   | <u>CGTGGCGCTT</u>  | <u>TCTCAATGCT</u> | 5500 |
| <u>CACGCTGTAG</u> | <u>GTATCTCACT</u>  | <u>TCGGTGTAGG</u>  | <u>TCGTTGGCTC</u>  | <u>CAAGCTGGGC</u> | 5550 |
| <u>TGTGTGCACG</u> | <u>AAACCCCGT</u>   | <u>TCAGCCCGAC</u>  | <u>CGCTGCGGCT</u>  | <u>TATCCGGTAA</u> | 5600 |
| <u>CTATCGTCTT</u> | <u>GAGTCCAACC</u>  | <u>CGGTAAGACA</u>  | <u>CGACITATCG</u>  | <u>CCACTGGCAG</u> | 5650 |
| <u>CAGCCACTGG</u> | <u>TAACAGGATT</u>  | <u>AGCAGAGCGA</u>  | <u>GGTATGTAGG</u>  | <u>CGGTGCTACA</u> | 5700 |

## FIGURE 6 (cont.)

| 10                 | 20                 | 30                 | 40                 | 50                |      |
|--------------------|--------------------|--------------------|--------------------|-------------------|------|
| <u>1234567890</u>  | <u>1234567890</u>  | <u>1234567890</u>  | <u>1234567890</u>  | <u>1234567890</u> |      |
| <u>GAGTTCTTGA</u>  | <u>AGTGGGIGGCC</u> | <u>TAACTAOGGC</u>  | <u>TACACTAGAA</u>  | <u>GGACAGTATT</u> | 5750 |
| <u>TGGTATCTGC</u>  | <u>GCTCTGCTGA</u>  | <u>AGCCAGTTAC</u>  | <u>CITCGGAAAAA</u> | <u>AGAGITGGTA</u> | 5800 |
| <u>GCTCTTIGATC</u> | <u>CGGCAAACAA</u>  | <u>ACCACCGCTG</u>  | <u>GTAGCGGIGG</u>  | <u>TTTTTTGTT</u>  | 5850 |
| <u>TGCAAGCAGC</u>  | <u>AGATTACCGG</u>  | <u>CAGAAAAAAA</u>  | <u>GGATCTCAAG</u>  | <u>AAGATCCTT</u>  | 5900 |
| <u>GATCTTTCT</u>   | <u>ACGGGGCTTG</u>  | <u>ACGCTCAGTG</u>  | <u>GAACGAAAAC</u>  | <u>TCACGTTAAG</u> | 5950 |
| <u>GGATTTTGGT</u>  | <u>CATGAGAGTA</u>  | <u>TCAAAAAGGA</u>  | <u>TCTTCACCTA</u>  | <u>GATCCTTTA</u>  | 6000 |
| <u>AATTAAAAAT</u>  | <u>GAAGTTTAA</u>   | <u>ATCAATCTAA</u>  | <u>AGTATATATG</u>  | <u>AGTAAACCTG</u> | 6050 |
| <u>GCTGACAGT</u>   | <u>TACCAATGCT</u>  | <u>TAATCAGTGA</u>  | <u>GGCACCTATC</u>  | <u>TCAGOGATCT</u> | 6100 |
| y1GelIreS          | ueL...siHo         | rPlaV...gr         | AueIreSgrA         |                   |      |
| <u>GTCTATTTCG</u>  | <u>TTCATCCATA</u>  | <u>GTTGCGTGAC</u>  | <u>TCCCCGTCGT</u>  | <u>GTAGATAACT</u> | 6150 |
| psAelIulGn         | sAteMprTue         | lnlGgrAlaV         | y1GgrAgrAr         | hTreSueL..        |      |
| <u>ACGATACGGG</u>  | <u>AGGGCTTACC</u>  | <u>ATCTGGCCCC</u>  | <u>AGTGCTGCAA</u>  | <u>TGATACCGCG</u> | 6200 |
| .reSlaVorP         | orPreSlaVt         | eMn1Gy1Gpr         | TsiHn1GueL         | reSlaValAu        |      |
| <u>AGACCCACGC</u>  | <u>TCACCGGCTC</u>  | <u>CAGATTATAC</u>  | <u>AGCAATAAAC</u>  | <u>CAGCCAGCG</u>  | 6250 |
| eLy1GlaVre         | SlaVorPulG         | ueLnsAelIu         | eLueLueLyl         | GalAueIgrA        |      |
| <u>GAAGGGCCGA</u>  | <u>GCGCAGAAGT</u>  | <u>GGTCCCTGCAA</u> | <u>CTTTATCCGC</u>  | <u>CICCATCCAG</u> | 6300 |
| ehPorPgrAa         | 1AsyCehPsi         | HpsAn1GueL         | syLelIgrAg         | rAprTylGrh        |      |
| <u>TCTATTAATT</u>  | <u>GTGCGGGGA</u>   | <u>AGCTAGAGTA</u>  | <u>AGTAGTTCGC</u>  | <u>CAGTTAATAG</u> | 6350 |
| T.....nsA          | nsAylGorPu         | eL...ueLue         | lryThsAalA         | ueL...ryTh        |      |
| <u>TTTGGCGAAC</u>  | <u>GTGCTGCGCA</u>  | <u>TIGCTACAGG</u>  | <u>CATOGTGGTG</u>  | <u>TCAOGCTCGT</u> | 6400 |
| sAalAsyOgr         | An1Gn1GprT         | nLG...ueLs         | yCgrAorPrh         | TlaVreSrhT        |      |
| <u>CGTTGGTAT</u>   | <u>GGCTTCATTTC</u> | <u>AGCTCOGGTT</u>  | <u>CCCAAOGATC</u>  | <u>AAGGCGAGTT</u> | 6450 |
| rhTh1GryTo         | rPsyLteM..         | .reSgrAnsA         | y1GlaVelIu         | eLalAueL..        |      |
| <u>ACATGATCCC</u>  | <u>CCATGGTGTG</u>  | <u>CAAAAAAGCG</u>  | <u>GTTAGCTCCT</u>  | <u>TCGGTCCTCC</u> | 6500 |
| .teMeliylG         | prTrhTrhts         | yCehPueLor         | P...reSgrA         | grApsAulGr        |      |
| <u>GATCGTTGTC</u>  | <u>AGAAGTAAGT</u>  | <u>TGGCGCGAGT</u>  | <u>GTTATCACTC</u>  | <u>ATGGTTATGG</u> | 6550 |
| eSgrAn1G...        | .ehPryTrhT         | orPgrAueLr         | hTel1Iav..         | .orP...orP        |      |
| <u>CAGCACTGCA</u>  | <u>TAATTCTCTT</u>  | <u>ACTGTCATGC</u>  | <u>CATCCGTAAG</u>  | <u>ATGCTTTCT</u>  | 6600 |
| ueLlaValAr         | yThsAulG..         | .nLG...alA         | teMgrAueLe         | lIreSsylnl        |      |
| <u>GTGACTGGTG</u>  | <u>AGTACTCAAC</u>  | <u>CAAGTCATTIC</u> | <u>TGAGAATAGT</u>  | <u>GTATGCGCG</u>  | 6650 |
| GreSnlGsiH         | rhTreSueLp         | rTrhTteMgr         | AueLeIrhT          | ryTalAalA1        |      |

FIGURE 6 (cont.)

| 10                 | 20                | 30                | 40                | 50                |      |
|--------------------|-------------------|-------------------|-------------------|-------------------|------|
| <u>1234567890</u>  | <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> | <u>1234567890</u> |      |
| ACCGAGITGC         | TCTTGCGCGG        | CGTCAATACT        | GGATAATACT        | GCGCCACATA        | 6700 |
| avReSnsAre         | SsyLylGorP        | rhTueLlavo        | rPryTryTgr        | AalAlaVryT        |      |
| GCAGAACTTT         | AAAAGTGCTC        | ATCATTGGAA        | AACGTTCTTC        | GGGGCGAAAA        | 6750 |
| syCehPsyLu         | eLuelLalA...      | ....n1GehP        | 1aVnsAsyLo        | rPalAehPla        |      |
| CTCTCAAGGA         | TCTTACCGCT        | GTTGAGATCC        | AGTTGATGT         | AAACCCACTCG       | 6800 |
| VgrAueLreS         | grAlaValAr        | hTreSelIpr        | TnsAreSrht        | 1aVprTulGs        |      |
| TGCACCCAAC         | TGATCTTCAG        | CATCTTTAC         | TTTCAACCAGC       | GTTCTGGGT         | 6850 |
| iH1aVprTre         | SelIsyLueL        | teMsyL...s        | yL...prTgr        | AsyLn1GrhT        |      |
| GAGCAAAAC          | AGGAAGGCAA        | AATGCGCAA         | AAAAGGGAAT        | AAGGGCGACA        | 6900 |
| ueLueLehPu         | eLehPalAeh        | PsiHgrAueL        | ehPorPehPu        | eLorPreSla        |      |
| CGGAAATGTT         | GAATACTCAT        | ACTCTCCCT         | TTTCAATATT        | ATTGAAGCAT        | 6950 |
| VreSelInsA         | ehPlav...1        |                   |                   |                   |      |
| TTATCAGGGT         | TATTGCTCA         | TGAGCGGATA        | CATATTGAA         | TGTATTTAGA        | 7000 |
| AAAATAAAC          | AATAGGGTT         | CGCGCACAT         | TTCCCCGAAA        | AGTGCCACCT        | 7050 |
| GAOGCTAAG          | AAACCATTAT        | TATCATGACA        | TTAACCTATA        | AAAATAGGCG        | 7100 |
| TATCAOGAGG         | CCCTTTOGTC        | TOGCGCGTT         | CGGTGATGAC        | GGTGAAAACC        | 7150 |
| TCTGACACAT         | GCAGCTCCG         | GAGACGGTCA        | CAGCTTGCT         | GTAAGCGGAT        | 7200 |
| CCCGGGAGCA         | GACAAGCGCG        | TCAGGGCGCG        | TCAGCGGGGIG       | TTGGCGGGIG        | 7250 |
| TCGGGGCTGG         | CTTAACATATG       | CGGCATCAGA        | GCAGATTGTA        | CTGAGAGTGC        | 7300 |
| ACCATATGCG         | GTGTGAAATA        | CGCACAGAT         | CGTAAGGAG         | AAAATACCGC        | 7350 |
| ATCAGGAAAT         | <u>TGTAAACGTT</u> | <u>AATATTGTT</u>  | <u>TAATTCGC</u>   | <u>GTAAATT</u>    | 7400 |
| <u>TGTTAAATCA</u>  | <u>GCTCATTTT</u>  | <u>TAACCAATAG</u> | <u>GGCGAAATCG</u> | <u>GCAAAATCCC</u> | 7450 |
| <u>TTATAAAATCA</u> | <u>AAAGAATAGA</u> | <u>CGAGATAGG</u>  | <u>GTGAGTGTT</u>  | <u>GTGCGAGTT</u>  | 7500 |
| <u>GGAACAAGAG</u>  | <u>TCCACTATTA</u> | <u>AAGAACGTGG</u> | <u>ACTCCAACGT</u> | <u>CAAAGGGCGA</u> | 7550 |
| <u>AAAACCGTCT</u>  | <u>ATCAGGGCGA</u> | <u>TGGCCACTA</u>  | <u>CGTGAACCAT</u> | <u>CACCTAATC</u>  | 7600 |

## FIGURE 6 (cont.)

| 10                | 20                | 30                 | 40                | 50                 |      |
|-------------------|-------------------|--------------------|-------------------|--------------------|------|
| 1234567890        | 1234567890        | 1234567890         | 1234567890        | 1234567890         |      |
| <u>AAGTTTTTIG</u> | <u>GGGTGAGGT</u>  | <u>GGCGTAAAGC</u>  | <u>ACTAAATCGG</u> | <u>AACCCCTAAAG</u> | 7650 |
| <u>GGAGGCCCCG</u> | <u>ATTTAGAGCT</u> | <u>TGACGGGAA</u>   | <u>AGCCGGGAA</u>  | <u>CGTGGGAGA</u>   | 7700 |
| <u>AAGGAAGGGA</u> | <u>AGAAAGCGAA</u> | <u>AGGAGGGGC</u>   | <u>GCTAGGGGCG</u> | <u>TGGCAAGTGT</u>  | 7750 |
| <u>AGCGGTACG</u>  | <u>CTGGGGTAA</u>  | <u>CCACCAACCC</u>  | <u>GGCCGGGCTT</u> | <u>AATGCGCGC</u>   | 7800 |
| <u>TACAGGGCGC</u> | <u>GTCGCGOCAT</u> | <u>TGCCCCATTCA</u> | <u>GGCTAOGCAA</u> | <u>CTGTTGGGAA</u>  | 7850 |
| <u>GGGCGATCGG</u> | <u>TGCGGGCCTC</u> | <u>TTCGCTATTAA</u> | <u>CGCCAGCTGG</u> | <u>CTGCAGGGGG</u>  | 7900 |
| <u>GGGGGGGGGG</u> | <u>GGGT</u>       |                    |                   |                    | 7914 |